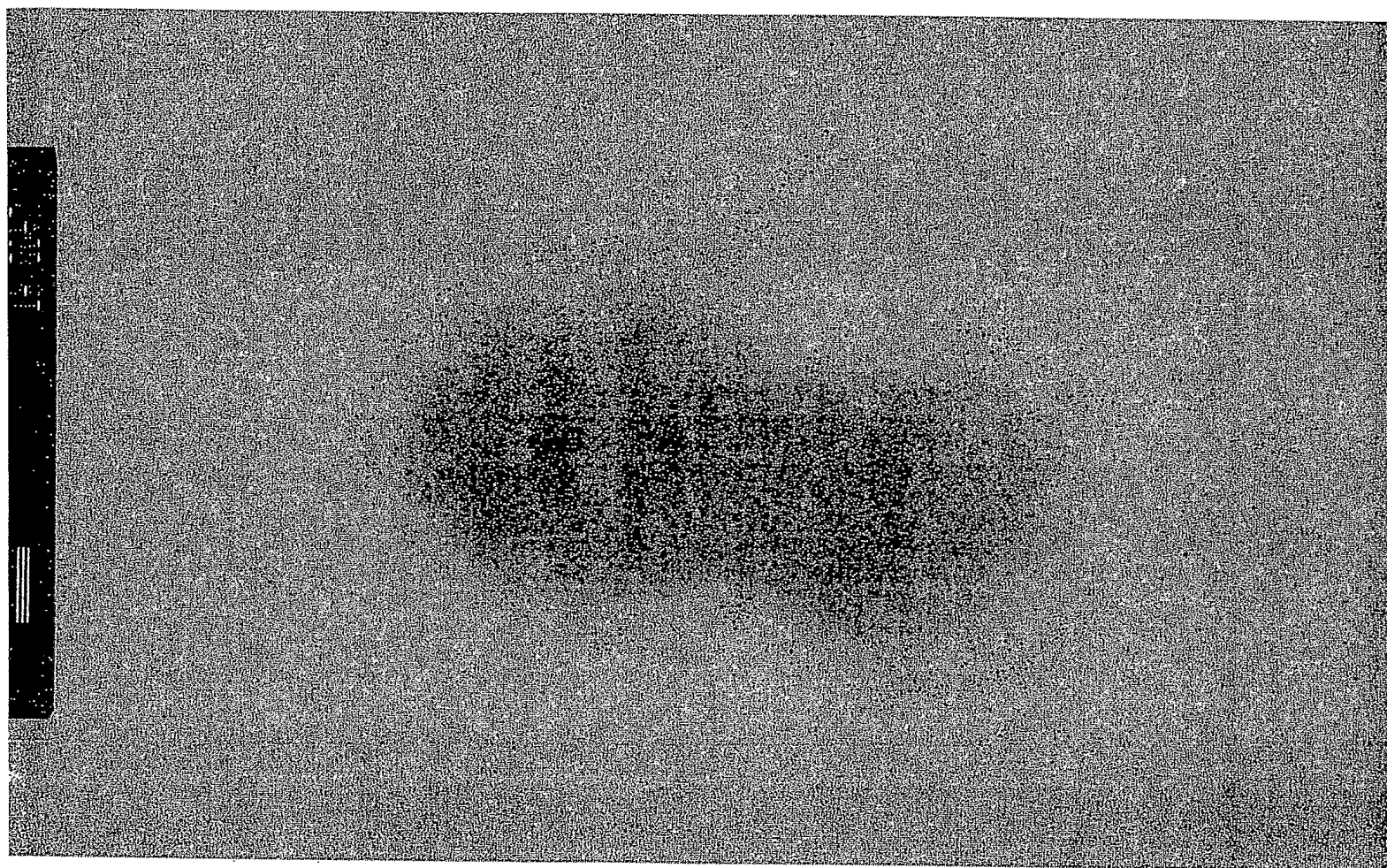


Figure 130



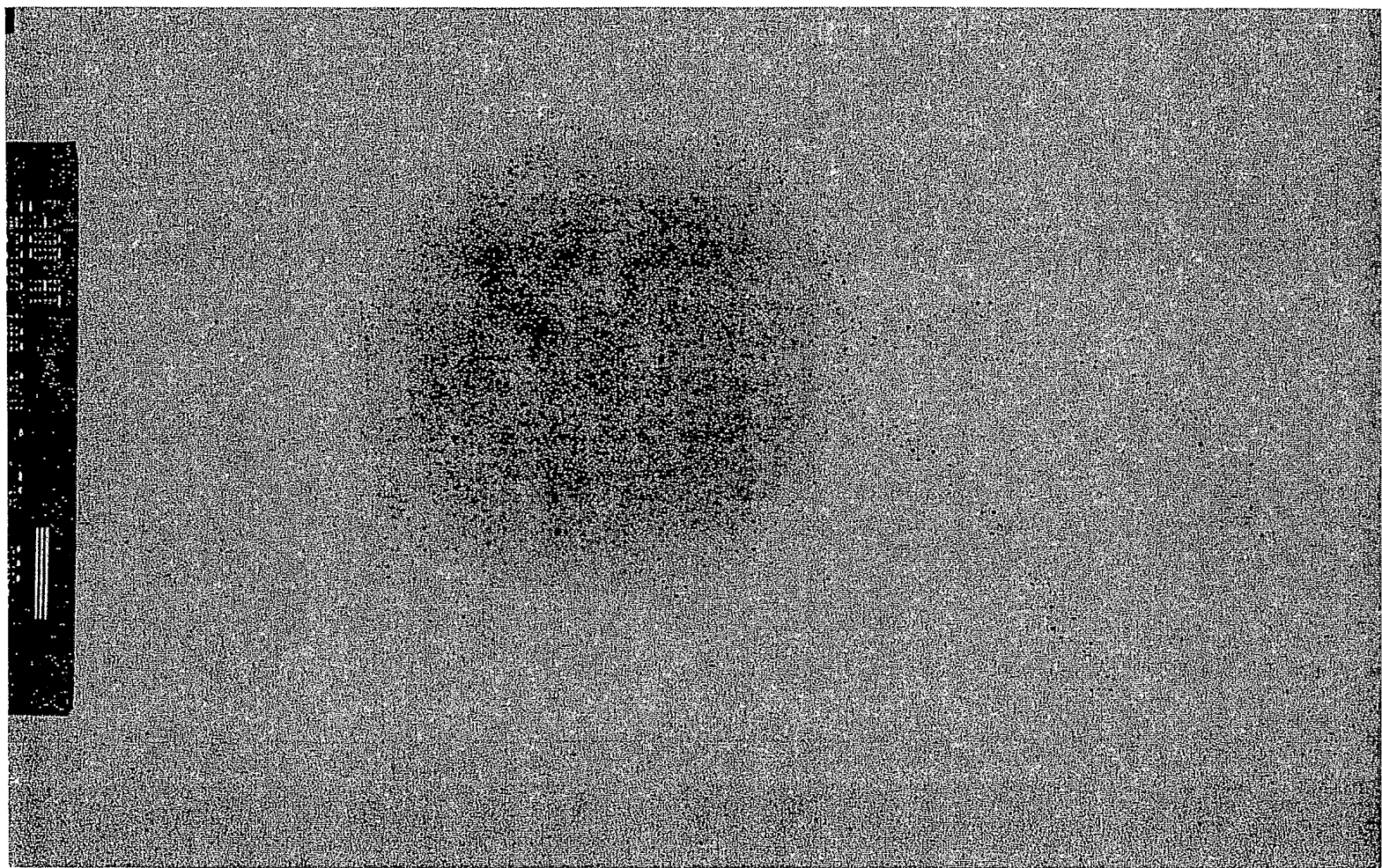
PCT/US05/27239 322/487

Figure 131



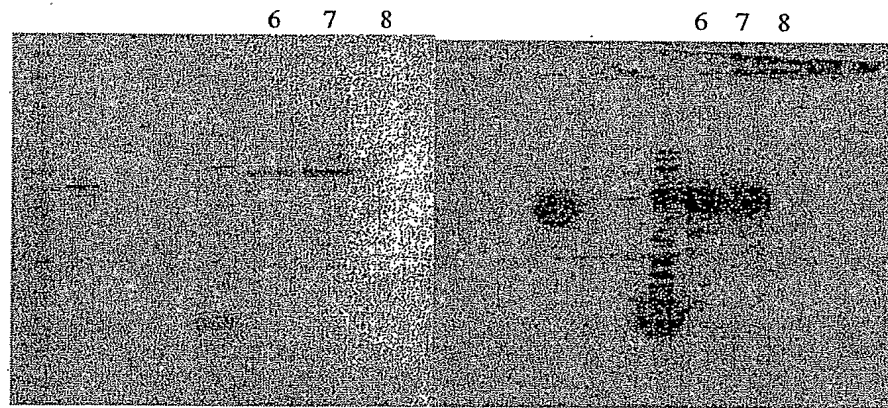
323/487

Figure 132



324/487

Figure 13.3

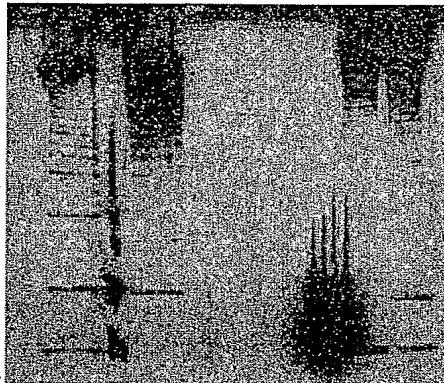


A

B

Figure 134

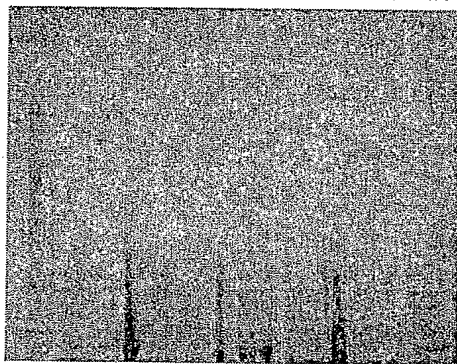
1 2 3 4 5 6 7 8 9 10



PCT/US05/27239 326/487

Figure 135

1 2 3 4 5 6 7 8 9 10



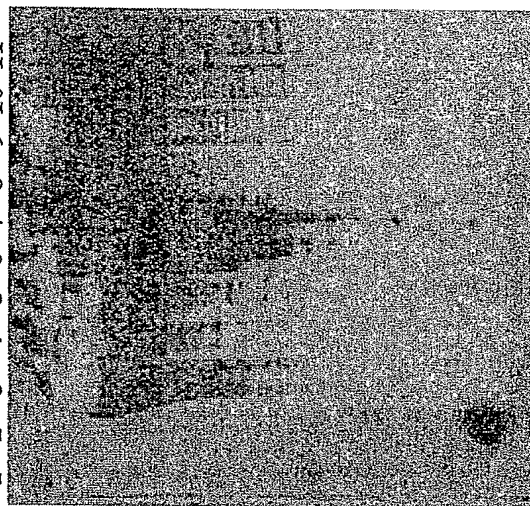
Pilus released by *Lactococcus* sonication

Figure 136A

1. MK
2. GBS 80 (10 ng)
3. L.lactis-A11 starting material (30', 0.2 OD)
4. L.lactis-A11 (not boiled, 0.33 OD)
5. L.lactis-A11 (5', 0.33 OD)
6. L.lactis-A11 (60' d, 0.33 OD)
7. L.lactis-A11 (30', 0.33 OD)
8. Supernatant (not boiled, 2 OD)
9. Supernatant (5', 2 OD)
10. Supernatant (30', 2 OD)
11. Supernatant (60', 2 OD)

SONICATED

starting material	pellet	supernatant
↓	↓	↓
1 2 3 4	5 6 7 8	9 10 11



α 80

Figure 136B

1. MK
2. L.lactis-A11
3. L.lactis-A11 starting material (30', 0.2 OD)
4. L.lactis-A11 (not boiled, 0.33 OD)
5. L.lactis-A11 (5', 0.33 OD)
6. L.lactis-A11 (60' d, 0.33 OD)
7. L.lactis-A11 (30', 0.33 OD)
8. Supernatant (not boiled, 2 OD)
9. Supernatant (5', 2 OD)
10. Supernatant (30', 2 OD)
11. Supernatant (60', 2 OD)

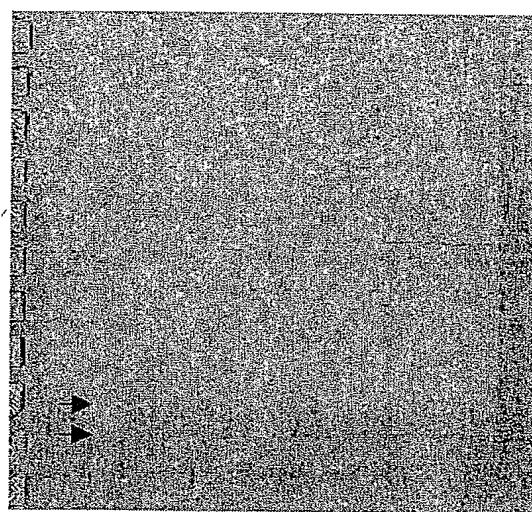
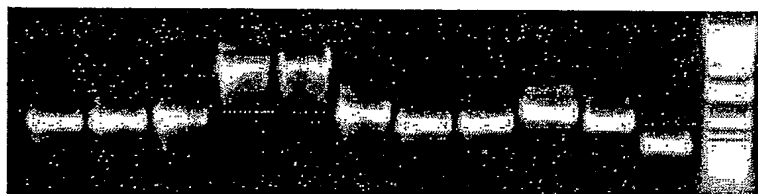




Figure 137



A



TIGR4

B

PCR product	contig_length _TIGR4	overlap
1	754	83
2	759	84
3	847	98
4	2550	99
5	2736	99
6	925	99
7	745	87
8	765	94
9	1008	94
10	802	64
11	461	

Figure 139



Figure 141A

ORF2_14CSR	MLNKYIEKRITDKITILNILLDIRSIELDELSTLTSLQSKSLLSILQELQETFEELTFN
ORF2_19AH	MLNKYIEKRITDKITILNILLDIRSIELDELSTLTSLQSKSLLSILQELQETFEELTFN
ORF2_19FTW	MLNKYIEKRITDKITILNILLDIRSIELDELSTLTSLQSKSLLSILQELQETFEELTFN
ORF2_23FP	MLNKYIEKRITDKITILNILLDIRSIELDELSTLTSLQSKSLLSILQELQETFEELTFN
ORF2_23FTW	MLNKYIEKRITDKITILNILLDIRSIELDELSTLTSLQSKSLLSILQELQETFEELTFN
ORF2_670	MLNKYIEKRITDKITILNILLDIRSIELDELSTLTSLQSKSLLSILQELQETFEELTFN
ORF2_6BF	MLNKYIEKRITDKITILNILLDIRSIELDELSTLTSLQSKSLLSILQELQETFEELTFN
ORF2_6BSP	MLNKYIEKRITDKITILNILLDIRSIELDELSTLTSLQSKSLLSILQELQETFEELTFN
ORF2_TIGR	MLNKYIEKRITDKITILNILLDIRSIELDELSTLTSLQSKSLLSILQELQETFEELTFN
ORF2_9VSP	MLNKYIEKRITDKITILNILLDIRSIELDELSTLTSLQSKSLLSILQELQETFEELTFN

ORF2_14CSR	LDTQQVQLIEHSHQTNYYFHQLYNQSTILKILRFFLLQGNQSFNEFTQKEYISIATGYR
ORF2_19AH	LDTQQVQLIEHSHQTNYYFHQLYNQSTILKILRFFLLQGNQSFNEFTQKEYISIATGYR
ORF2_19FTW	LDTQQVQLIEHSHQTNYYFHQLYNQSTILKILRFFLLQGNQSFNEFTQKEYISIATGYR
ORF2_23FP	LDTQQVQLIEHSHQTNYYFHQLYNQSTILKILRFFLLQGNQSFNEFTQKEYISIATGYR
ORF2_23FTW	LDTQQVQLIEHSHQTNYYFHQLYNQSTILKILRFFLLQGNQSFNEFTQKEYISIATGYR
ORF2_670	LDTQQVQLIEHSHQTNYYFHQLYNQSTILKILRFFLLQGNQSFNEFTQKEYISIATGYR
ORF2_6BF	LDTQQVQLIEHSHQTNYYFHQLYNQSTILKILRFFLLQGNQSFNEFTQKEYISIATGYR
ORF2_6BSP	LDTQQVQLIEHSHQTNYYFHQLYNQSTILKILRFFLLQGNQSFNEFTQKEYISIATGYR
ORF2_TIGR	LDTQQVQLIEHSHQTNYYFHQLYNQSTILKILRFFLLQGNQSFNEFTQKEYISIATGYR
ORF2_9VSP	LDTQQVQLIEHSHQTNYYFHQLYNQSTILKILRFFLLQGNQSFNEFTQKEYISIATGYR

ORF2_14CSR	VRQKCGLLRSVGLDLVKNQVVGPEYRIRFLIALLOFHFGIEIYDLNDGSMWVTHMIVQ
ORF2_19AH	VRQKCGLLRSVGLDLVKNQVVGPEYRIRFLIALLOFHFGIEIYDLNDGSMWVTHMIVQ
ORF2_19FTW	VRQKCGLLRSVGLDLVKNQVVGPEYRIRFLIALLOFHFGIEIYDLNDGSMWVTHMIVQ
ORF2_23FP	VRQKCGLLRSVGLDLVKNQVVGPEYRIRFLIALLOFHFGIEIYDLNDGSMWVTHMIVQ
ORF2_23FTW	VRQKCGLLRSVGLDLVKNQVVGPEYRIRFLIALLOFHFGIEIYDLNDGSMWVTHMIVQ
ORF2_670	VRQKCGLLRSVGLDLVKNQVVGPEYRIRFLIALLOFHFGIEIYDLNDGSMWVTHMIVQ
ORF2_6BF	VRQKCGLLRSVGLDLVKNQVVGPEYRIRFLIALLOFHFGIEIYDLNDGSMWVTHMIVQ
ORF2_6BSP	VRQKCGLLRSVGLDLVKNQVVGPEYRIRFLIALLOFHFGIEIYDLNDGSMWVTHMIVQ
ORF2_TIGR	VRQKCGLLRSVGLDLVKNQVVGPEYRIRFLIALLOFHFGIEIYDLNDGSMWVTHMIVQ
ORF2_9VSP	VRQKCGLLRSVGLDLVKNQVVGPEYRIRFLIALLOFHFGIEIYDLNDGSMWVTHMIVQ

ORF2_14CSR	SNSQLSHELLEITPDEYVHFSILVALTWKRREFPLEFPESKEFEKLKNLFMPILMEHCQ
ORF2_19AH	SNSQLSHELLEITPDEYVHFSILVALTWKRREFPLEFPESKEFEKLKNLFMPILMEHCQ
ORF2_19FTW	SNSQLSHELLEITPDEYVHFSILVALTWKRREFPLEFPESKEFEKLKNLFMPILMEHCQ
ORF2_23FP	SNSQLSHELLEITPDEYVHFSILVALTWKRREFPLEFPESKEFEKLKNLFMPILMEHCQ
ORF2_23FTW	SNSQLSHELLEITPDEYVHFSILVALTWKRREFPLEFPESKEFEKLKNLFMPILMEHCQ
ORF2_670	SNSQLSHELLEITPDEYVHFSILVALTWKRREFPLEFPESKEFEKLKNLFMPILMEHCQ
ORF2_6BF	SNSQLSHELLEITPDEYVHFSILVALTWKRREFPLEFPESKEFEKLKNLFMPILMEHCQ
ORF2_6BSP	SNSQLSHELLEITPDEYVHFSILVALTWKRREFPLEFPESKEFEKLKNLFMPILMEHCQ
ORF2_TIGR	SNSQLSHELLEITPDEYVHFSILVALTWKRREFPLEFPESKEFEKLKNLFMPILMEHCQ
ORF2_9VSP	SNSQLSHELLEITPDEYVHFSILVALTWKRREFPLEFPESKEFEKLKNLFMPILMEHCQ

ORF2_14CSR	TYLEPHANMTFTQEELDYIFLVYCSANSSFSKDKWNQEKKTHTIQLILQHTRGKHLSSKF
ORF2_19AH	TYLEPHANMTFTQEELDYIFLVYCSANSSFSKDKWNQEKKTHTIQLILQHTRGKHLSSKF
ORF2_19FTW	TYLEPHANMTFTQEELDYIFLVYCSANSSFSKDKWNQEKKTHTIQLILQHTRGKHLSSKF
ORF2_23FP	TYLEPHANMTFTQEELDYIFLVYCSANSSFSKDKWNQEKKTHTIQLILQHTRGKHLSSKF
ORF2_23FTW	TYLEPHANMTFTQEELDYIFLVYCSANSSFSKDKWNQEKKTHTIQLILQHTRGKHLSSKF
ORF2_670	TYLEPHANMTFTQEELDYIFLVYCSANSSFSKDKWNQEKKTHTIQLILQHTRGKHLSSKF
ORF2_6BF	TYLEPHANMTFTQEELDYIFLVYCSANSSFSKDKWNQEKKTHTIQLILQHTRGKHLSSKF
ORF2_6BSP	TYLEPHANMTFTQEELDYIFLVYCSANSSFSKDKWNQEKKTHTIQLILQHTRGKHLSSKF
ORF2_TIGR	TYLEPHANMTFTQEELDYIFLVYCSANSSFSKDKWNQEKKTHTIQLILQHTRGKHLSSKF
ORF2_9VSP	TYLEPHANMTFTQEELDYIFLVYCSANSSFSKDKWNQEKKTHTIQLILQHTRGKHLSSKF

Figure 141B

ORF2_14CSR KNILGNDISNSLSFLTALTFLTRTFLFGLQNLVPYNNYEHYGIIESDKPLYHISKAIVQE
ORF2_19AH KNILGNDISNSLSFLTALTFLTRTFLFGLQNLVPYNNYEHYGIIESDKPLYHISKAIVQE
ORF2_19FTW KNILGNDISNSLSFLTALTFLTRTFLFGLQNLVPYNNYEHYGIIESDKPLYHISKAIVQE
ORF2_23FP KNILGNDISNSLSFLTALTFLTRTFLFGLQNLVPYNNYEHYGIIESDKPLYHISKAIVQE
ORF2_23FTW KNILGNDISNSLSFLTALTFLTRTFLFGLQNLVPYNNYEHYGIIESDKPLYHISKAIVQE
ORF2_670 KNILGNDISNSLSFLTALTFLTRTFLFGLQNLVPYNNYEHYGIIESDKPLYHISKAIVQE
ORF2_6BF KNILGNDISNSLSFLTALTFLTRTFLFGLQNLVPYNNYEHYGIIESDKPLYHISKAIVQE
ORF2_6BSP KNILGNDISNSLSFLTALTFLTRTFLFGLQNLVPYNNYEHYGIIESDKPLYHISKAIVQE
ORF2_TIGR KNILGNDISNSLSFLTALTFLTRTFLFGLQNLVPYNNYEHYGIIESDKPLYHISKAIVQE
ORF2_9VSP KNILGNDISNSLSFLTALTFLTRTFLFGLQNLVPYNNYEHYGIIESDKPLYHISKAIVQE

ORF2_14CSR WMTEQKIEGVIDQHRLYLFSLYLTETIFSSSLPAIPIFIILNNQADVNLIKSIILRNFTDK
ORF2_19AH WMTEQKIEGVIDQHRLYLFSLYLTETIFSSSLPAIPIFIILNNQADVNLIKSIILRNFTDK
ORF2_19FTW WMTEQKIEGVIDQHRLYLFSLYLTETIFSSSLPAIPIFIILNNQADVNLIKSIILRNFTDK
ORF2_23FP WMTEQKIEGVIDQHRLYLFSLYLTETIFSSSLPAIPIFIILNNQADVNLIKSIILRNFTDK
ORF2_23FTW WMTEQKIEGVIDQHRLYLFSLYLTETIFSSSLPAIPIFIILNNQADVNLIKSIILRNFTDK
ORF2_670 WMTEQKIEGVIDQHRLYLFSLYLTETIFSSSLPAIPIFIILNNQADVNLIKSIILRNFTDK
ORF2_6BF WMTEQKIEGVIDQHRLYLFSLYLTETIFSSSLPAIPIFIILNNQADVNLIKSIILRNFTDK
ORF2_6BSP WMTEQKIEGVIDQHRLYLFSLYLTETIFSSSLPAIPIFIILNNQADVNLIKSIILRNFTDK
ORF2_TIGR WMTEQKIEGVIDQHRLYLFSLYLTETIFSSSLPAIPIFIILNNQADVNLIKSIILRNFTDK
ORF2_9VSP WMTEQKIEGVIDQHRLYLFSLYLTETIFSSSLPAIPIFIILNNQADVNLIKSIILRNFTDK

ORF2_14CSR VASVTGYNILISPPPSEHLTEPLIIITTKKEYLPYVKKQYPKGKHHFLTIALDLHVSQQR
ORF2_19AH VASVTGYNILISPPPSEHLTEPLIIITTKKEYLPYVKKQYPKGKHHFLTIALDLHVSQQR
ORF2_19FTW VASVTGYNILISPPPSEHLTEPLIIITTKKEYLPYVKKQYPKGKHHFLTIALDLHVSQQR
ORF2_23FP VASVTGYNILISPPPSEHLTEPLIIITTKKEYLPYVKKQYPKGKHHFLTIALDLHVSQQR
ORF2_23FTW VASVTGYNILISPPPSEHLTEPLIIITTKKEYLPYVKKQYPKGKHHFLTIALDLHVSQQR
ORF2_670 VASVTGYNILISPPPSEHLTEPLIIITTKKEYLPYVKKQYPKGKHHFLTIALDLHVSQQR
ORF2_6BF VASVTGYNILISPPPSEHLTEPLIIITTKKEYLPYVKKQYPKGKHHFLTIALDLHVSQQR
ORF2_6BSP VASVTGYNILISPPPSEHLTEPLIIITTKKEYLPYVKKQYPKGKHHFLTIALDLHVSQQR
ORF2_TIGR VASVTGYNILISPPPSEHLTEPLIIITTKKEYLPYVKKQYPKGKHHFLTIALDLHVSQQR
ORF2_9VSP VASVTGYNILISPPPSEHLTEPLIIITTKKEYLPYVKKQYPKGKHHFLTIALDLHVSQQR

ORF2_14CSR LIYQTIVDIRKEAFDKRVAMIAKKAHYLL
ORF2_19AH LIYQTIVDIRKEAFDKRVAMIAKKAHYLL
ORF2_19FTW LIYQTIVDIRKEAFDKRVAMIAKKAHYLL
ORF2_23FP LIYQTIVDIRKEAFDKRVAMIAKKAHYLL
ORF2_23FTW LIYQTIVDIRKEAFDKRVAMIAKKAHYLL
ORF2_670 LIYQTIVDIRKEAFDKRVAMIAKKAHYLL
ORF2_6BF LIYQTIVDIRKEAFDKRVAMIAKKAHYLL
ORF2_6BSP LIYQTIVDIRKEAFDKRVAMIAKKAHYLL
ORF2_TIGR LIYQTIVDIRKEAFDKRVAMIAKKAHYLL
ORF2_9VSP LIYQTIVDIRKEAFDKRVAMIAKKAHYLL

Figure 142A

```
ORF3_19AH      MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVF
ORF3_23FP      MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVF
ORF3_14CSR     MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVF
ORF3_670       MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVF
ORF3_6BF       MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVF
ORF3_6BSP      MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVF
ORF3_19FTW     MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVF
ORF3_9VSP      MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVF
ORF3_23FTW     MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVF
ORF3_TIGR      MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSIPAIGKVVIKETGEGGALLGDAVF
*****

ORF3_19AH      ELKNNTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQP PVGYKPSTKQWTVVEVEKNGRT
ORF3_23FP      ELKNNTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQP PVGYKPSTKQWTVVEVEKNGRT
ORF3_14CSR     ELKNNTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQP PVGYKPSTKQWTVVEVEKNGRT
ORF3_670       ELKNNTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQP PVGYKPSTKQWTVVEVEKNGRT
ORF3_6BF       ELKNNTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQP PVGYKPSTKQWTVVEVEKNGRT
ORF3_6BSP      ELKNNTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQP PVGYKPSTKQWTVVEVEKNGRT
ORF3_19FTW     ELKNNTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQP PVGYKPSTKQWTVVEVEKNGRT
ORF3_9VSP      ELKNNTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQP PVGYKPSTKQWTVVEVEKNGRT
ORF3_23FTW     ELKNNTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQP PVGYKPSTKQWTVVEVEKNGRT
ORF3_TIGR      ELKNNTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQP PVGYKPSTKQWTVVEVEKNGRT
*****

ORF3_19AH      TVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVIPEG
ORF3_23FP      TVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVIPEG
ORF3_14CSR     TVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVIPEG
ORF3_670       TVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVIPEG
ORF3_6BF       TVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVIPEG
ORF3_6BSP      TVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVIPEG
ORF3_19FTW     TVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVIPEG
ORF3_9VSP      TVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVIPEG
ORF3_23FTW     TVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVIPEG
ORF3_TIGR      TVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVIPEG
*****

ORF3_19AH      TLSKRIYQVNNLDDNQYGIELTVSGKTTVETKEASTPLDVVILLDNSNSMSNIRHNHAHR
ORF3_23FP      TLSKRIYQVNNLDDNQYGIELTVSGKTTVETKEASTPLDVVILLDNSNSMSNIRHNHAHR
ORF3_14CSR     TLSKRIYQVNNLDDNQYGIELTVSGKTTVETKEASTPLDVVILLDNSNSMSNIRHNHAHR
ORF3_670       TLSKRIYQVNNLDDNQYGIELTVSGKTTVETKEASTPLDVVILLDNSNSMSNIRHNHAHR
ORF3_6BF       TLSKRIYQVNNLDDNQYGIELTVSGKTTVETKEASTPLDVVILLDNSNSMSNIRHNHAHR
ORF3_6BSP      TLSKRIYQVNNLDDNQYGIELTVSGKTTVETKEASTPLDVVILLDNSNSMSNIRHNHAHR
ORF3_19FTW     TLSKRIYQVNNLDDNQYGIELTVSGKTVYERKDKSVPLDVVILLDNSNSMSNIRKNARR
ORF3_9VSP      TLSKRIYQVNNLDDNQYGIELTVSGKTVYERKDKSVPLDVVILLDNSNSMSNIRKNARR
ORF3_23FTW     TLSKRIYQVNNLDDNQYGIELTVSGKTVYEQKDKSVPLDVVILLDNSNSMSNIRKNARR
ORF3_TIGR      TLSKRIYQVNNLDDNQYGIELTVSGKTVYEQKDKSVPLDVVILLDNSNSMSNIRKNARR
*****

ORF3_19AH      AEKAGEATRALVDKITSNPDNRVALVTYGSTIFDGSEATVEKGVADANGKILNDSALWTF
ORF3_23FP      AEKAGEATRALVDKITSNPDNRVALVTYGSTIFDGSEATVEKGVADANGKILNDSALWTF
ORF3_14CSR     AEKAGEATRALVDKITSNPDNRVALVTYGSTIFDGSEATVEKGVADANGKILNDSALWTF
ORF3_670       AEKAGEATRALVDKITSNPDNRVALVTYGSTIFDGSEATVEKGVADANGKILNDSALWTF
ORF3_6BF       AEKAGEATRALVDKITSNPDNRVALVTYGSTIFDGSEATVEKGVADANGKILNDSALWTF
ORF3_6BSP      AEKAGEATRALVDKITSNPDNRVALVTYGSTIFDGSEATVEKGVADANGKILNDSALWTF
ORF3_19FTW     AERAGEATRSIDKITSDPENRVALVTYASTIFDGTEFTVEKGVADKNGKRLNDSLEFNNY
ORF3_9VSP      AERAGEATRSIDKITSDPENRVALVTYASTIFDGTEFTVEKGVADKNGKRLNDSLEFNNY
ORF3_23FTW     AERAGEATRSIDKITSDPENRVALVTYASTIFDGTEFTVEKGVADKNGKRLNDSLEFNNY
ORF3_TIGR      AERAGEATRSIDKITSSENPRVALVTYASTIFDGTEFTVEKGVADKNGKRLNDSLEFNNY
**::*****::*****::*****::*****::*****::*****::*****::**::
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 DRTTFTAKTYNYSFLNLTSDPDTDIQTIKDRIPSDAEELNKDKLMYQFGATFTQKALMTAD
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 DQTSFTTNTKDYSLKLTNDKNDIVELKNKVPTEAEDHDGNRLMYQFGATFTQKALMKAD
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 DQTSFTTNTKDYSLKLTNDKNDIVELKNKVPTEAEDHDGNRLMYQFGATFTQKALMKAD
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DILTKQARPNSKKVIFHITDGVPTMSYPINFKYTGTTQSYRTQLNNFKAKTPNSSGILLE
DILTKQARPNSKKVIFHITDGVPTMSYPINFKYTGTTQSYRTQLNNFKAKTPNSSGILLE
DILTKQARPNSKKVIFHITDGVPTMSYPINFKYTGTTQSYRTQLNNFKAKTPNSSGILLE
DILTKQARPNSKKVIFHITDGVPTMSYPINFKYTGTTQSYRTQLNNFKAKTPNSSGILLE
DILTKQARPNSKKVIFHITDGVPTMSYPINFKYTGTTQSYRTQLNNFKAKTPNSSGILLE
EILTQQARQNSQKVI F H I T D G V P T M S Y P I N F N H A T F A P S Y Q N Q L N A F F S K S P N K D G I L L S
EILTQQARQNSQKVI F H I T D G V P T M S Y P I N F N H A T F A P S Y Q N Q L N V F F S K S P N K D G I L L S
EILTQQARQNSQKVI F H I T D G V P T M S Y P I N F N H A T F A P S Y Q N Q L N A F F S K S P N K D G I L L S
EILTQQARQNSQKVI F H I T D G V P T M S Y P I N F N H A T F A P S Y Q N Q L N A F F S K S P N K D G I L L S
: *:*****::: :*:***:~::~:~::~:

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DFV TWSADGEHKIVRGDGESYQMFTKKPVTDQYGVHQILSITSMEQRAKLV SAGYRFYGT
DFV TWSADGEHKIVRGDGESYQMFTKKPVTDQYGVHQILSITSMEQRAKLV SAGYRFYGT
DFV TWSADGEHKIVRGDGESYQMFTKKPVTDQYGVHQILSITSMEQRAKLV SAGYRFYGT
DFV TWSADGEHKIVRGDGESYQMFTKKPVTDQYGVHQILSITSMEQRAKLV SAGYRFYGT
DFV TWSADGEHKIVRGDGESYQMFTKKPVTDQYGVHQILSITSMEQRAKLV SAGYRFYGT
DFV TWSADGEHKIVRGDGESYQMFTKKPVTDQYGVHQILSITSMEQRAKLV SAGYRFYGT
DFITQATSGEHTIVRGDGQSYQMFTDKTVYEK-GAPAAFPVK-PEKYS E MKA VGYAVIGD
DFITQATSGEHTIVRGDGQSYQMFTDKTVYEK-GAPAAFPVK-PEKYS E MKA VGYAVIGD
DFITQATSGEHTIVRGDGQSYQMFTDKTVYEK-GAPAAFPVK-PEKYS E MKA VGYAVIGD
DFITQATSGEHTIVRGDGQSYQMFTDKTVYEK-GAPAAFPVK-PEKYS E MKA VGYAVIGD
**:* :.:***.*****:*****.*:* :.:* :.:* :.:* :.:* :.:* :.:*

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-----DLYLYWRDSILAYPFNSSTDWITNHGDPPTTWYYNGNMAQDGYDVFTVGVGVNGDP
-----DLYLYWRDSILAYPFNSSTDWITNHGDPPTTWYYNGNMAQDGYDVFTVGVGVNGDP
-----DLYLYWRDSILAYPFNSSTDWITNHGDPPTTWYYNGNMAQDGYDVFTVGVGVNGDP
-----DLYLYWRDSILAYPFNSSTDWITNHGDPPTTWYYNGNMAQDGYDVFTVGVGVNGDP
-----DLYLYWRDSILAYPFNSSTDWITNHGDPPTTWYYNGNMAQDGYDVFTVGVGVNGDP
-----DLYLYWRDSILAYPFNSSTDWITNHGDPPTTWYYNGNMAQDGYDVFTVGVGVNGDP
PINGGYIWLNWRESILAYPFNSNTAKITNHGAPTRWYYNGNIAPDGYDVFTVGVGIGINGDP
PINGGYIWLNWRESILAYPFNSNTAKITNHGDPTRWYYNGNIAPDGYDVFTVGVGIGINGDP
PINGGYIWLNWRESILAYPFNSNTAKITNHGDPTRWYYNGNIAPDGYDVFTVGVGIGINGDP
PINGGYIWLNWRESILAYPFNSNTAKITNHGDPTRWYYNGNIAPDGYDVFTVGVGIGINGDP
:.* **:*.....* ***** ** *****: *****.*.....*

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[illegible]

Figure 142C

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ORF3_19AH      IDFLQADGRFDPADYTLTANDGSSLVNNVPTGGPQNDGGLLKNKAVFYDTEKIRIVTG
ORF3_23FP      IDFLQADGRFDPADYTLTANDGSSLVNNVPTGGPQNDGGLLKNKAVFYDTEKIRIVTG
ORF3_14CSR     IDFLQADGRFDPADYTLTANDGSSLVNNVPTGGPQNDGGLLKNKAVFYDTEKIRIVTG
ORF3_670       IDFLQADGRFDPADYTLTANDGSSLVNNVPTGGPQNDGGLLKNKAVFYDTEKIRIVTG
ORF3_6BF       IDFLQADGRFDPADYTLTANDGSSLVNNVPTGGPQNDGGLLKNKAVFYDTEKIRIVTG
ORF3_6BSP      IDFLQADGRFDPADYTLTANDGSSLVNNVPTGGPQNDGGLLKNKAVFYDTEKIRIVTG
ORF3_19FTW     IDLQLGTDGRFDPADYTLTANDGSRLENGQAVGGPQNDGGLLKNKAVFYDTEKIRIVTG
ORF3_9VSP      IDLQLGTDGRFDPADYTLTANDGSRLENGQAVGGPQNDGGLLKNKAVFYDTEKIRIVTG
ORF3_23FTW     IDLQLGTDGRFDPADYTLTANDGSRLENGQAVGGPQNDGGLLKNKAVFYDTEKIRIVTG
ORF3_TIGR      IDLQLGTDGRFDPADYTLTANDGSRLENGQAVGGPQNDGGLLKNKAVFYDTEKIRIVTG
                ****.****.*****.*****.*****.*****.*****.*****.*****

ORF3_19AH      LYLGTGEKVTLTYNVRLNDQFVSNKFYDTNGRTTLHPKEVEKNTVRDFPIPKIRDVRKYP
ORF3_23FP      LYLGTGEKVTLTYNVRLNDQFVSNKFYDTNGRTTLHPKEVEKNTVRDFPIPKIRDVRKYP
ORF3_14CSR     LYLGTGEKVTLTYNVRLNDQFVSNKFYDTNGRTTLHPKEVEKNTVRDFPIPKIRDVRKYP
ORF3_670       LYLGTGEKVTLTYNVRLNDQFVSNKFYDTNGRTTLHPKEVEKNTVRDFPIPKIRDVRKYP
ORF3_6BF       LYLGTGEKVTLTYNVRLNDQFVSNKFYDTNGRTTLHPKEVEKNTVRDFPIPKIRDVRKYP
ORF3_6BSP      LYLGTGEKVTLTYNVRLNDQFVSNKFYDTNGRTTLHPKEVEKNTVRDFPIPKIRDVRKYP
ORF3_19FTW     LYLGTGEKVTLTYNVRLNDQFVSNKFYDTNGRTTLHPKEVEKNTVRDFPIPKIRDVRKYP
ORF3_9VSP      LYLGTGEKVTLTYNVRLNDQFVSNKFYDTNGRTTLHPKEVEKNTVRDFPIPKIRDVRKYP
ORF3_23FTW     LYLGTGEKVTLTYNVRLNDQFVSNKFYDTNGRTTLHPKEVEKNTVRDFPIPKIRDVRKYP
ORF3_TIGR      LYLGTGEKVTLTYNVRLNDQFVSNKFYDTNGRTTLHPKEVEKNTVRDFPIPKIRDVRKYP
                *****.*****.*****.*****.*****.*****.*****.*****

ORF3_19AH      EITIPKEKKLGEIEFIKINKNDKKPLRDAVFSLQKQHPDYDPIYGAIDQNGTYQNVRTGE
ORF3_23FP      EITIPKEKKLGEIEFIKINKNDKKPLRDAVFSLQKQHPDYDPIYGAIDQNGTYQNVRTGE
ORF3_14CSR     EITIPKEKKLGEIEFIKINKNDKKPLRDAVFSLQKQHPDYDPIYGAIDQNGTYQNVRTGE
ORF3_670       EITIPKEKKLGEIEFIKINKNDKKPLRDAVFSLQKQHPDYDPIYGAIDQNGTYQNVRTGE
ORF3_6BF       EITIPKEKKLGEIEFIKINKNDKKPLRDAVFSLQKQHPDYDPIYGAIDQNGTYQNVRTGE
ORF3_6BSP      EITIPKEKKLGEIEFIKINKNDKKPLRDAVFSLQKQHPDYDPIYGAIDQNGTYQNVRTGE
ORF3_19FTW     AITIAKEKKLGEIEFIKINKNDKKPLRDAVFSLQKQHPDYDPIYGAIDQNGTYQNVRTGE
ORF3_9VSP      AITIAKEKKLGEIEFIKINKNDKKPLRDAVFSLQKQHPDYDPIYGAIDQNGTYQNVRTGE
ORF3_23FTW     EITISKEKKLGDIEFIKVNKNDKKPLRDAVFSLQKQHPDYDPIYGAIDQNGTYQNVRTGE
ORF3_TIGR      EITISKEKKLGDIEFIKVNKNDKKPLRDAVFSLQKQHPDYDPIYGAIDQNGTYQNVRTGE
                *****.*****.*****.*****.*****.*****.*****.*****

ORF3_19AH      DGKLTFFKNLSDGKRIEENSEPAKPVQNKPIVAFQIVNGEVRDVTISIVPDIPAGYEF
ORF3_23FP      DGKLTFFKNLSDGKRIEENSEPAKPVQNKPIVAFQIVNGEVRDVTISIVPDIPAGYEF
ORF3_14CSR     DGKLTFFKNLSDGKRIEENSEPAKPVQNKPIVAFQIVNGEVRDVTISIVPDIPAGYEF
ORF3_670       DGKLTFFKNLSDGKRIEENSEPAKPVQNKPIVAFQIVNGEVRDVTISIVPDIPAGYEF
ORF3_6BF       DGKLTFFKNLSDGKRIEENSEPAKPVQNKPIVAFQIVNGEVRDVTISIVPDIPAGYEF
ORF3_6BSP      DGKLTFFKNLSDGKRIEENSEPAKPVQNKPIVAFQIVNGEVRDVTISIVPDIPAGYEF
ORF3_19FTW     DGKLTFFKNLSDGKRIEENSEPAKPVQNKPIVAFQIVNGEVRDVTISIVPDIPAGYEF
ORF3_9VSP      DGKLTFFKNLSDGKRIEENSEPAKPVQNKPIVAFQIVNGEVRDVTISIVPDIPAGYEF
ORF3_23FTW     DGKLTFFKNLSDGKRIEENSEPAKPVQNKPIVAFQIVNGEVRDVTISIVPDIPAGYEF
ORF3_TIGR      DGKLTFFKNLSDGKRIEENSEPAKPVQNKPIVAFQIVNGEVRDVTISIVPDIPAGYEF
                *****.*****.*****.*****.*****.*****.*****.*****

ORF3_19AH      TNDKHYITNEPIPPKREYPTGGIGMLPFYLGICMMMGGVLLYTRKNP
ORF3_23FP      TNDKHYITNEPIPPKREYPTGGIGMLPFYLGICMMMGGVLLYTRKNP
ORF3_14CSR     TNDKHYITNEPIPPKREYPTGGIGMLPFYLGICMMMGGVLLYTRKHP
ORF3_670       TNDKHYITNEPIPPKREYPTGGIGMLPFYLGICMMMGGVLLYTRKHP
ORF3_6BF       TNDKHYITNEPIPPKREYPTGGIGMLPFYLGICMMMGGVLLYTRKHP
ORF3_6BSP      TNDKHYITNEPIPPKREYPTGGIGMLPFYLGICMMMGGVLLYTRKHP
ORF3_19FTW     TNDKHYITNEPIPPKREYPTGGIGMLPFYLGICMMMGGVLLYTRKHP
ORF3_9VSP      TNDKHYITNEPIPPKREYPTGGIGMLPFYLGICMMMGGVLLYTRKHP
ORF3_23FTW     TNDKHYITNEPIPPKREYPTGGIGMLPFYLGICMMMGGVLLYTRKHP
ORF3_TIGR      TNDKHYITNEPIPPKREYPTGGIGMLPFYLGICMMMGGVLLYTRKHP
                *****.*****.*****.*****.*****.*****.*****.*****
```


Figure 143A

ORF4_6BF MKSINKFLTMLAALLLTASSLSFSAATVFAADNVSTAPDAVTKTLTIHKLLLSEDDLKTWD
 ORF4_6BSP MKSINKFLTMLAALLLTASSLSFSAATVFAADNVSTAPDAVTKTLTIHKLLLSEDDLKTWD
 ORF4_670 MKSINKFLTMLAALLLTASSLSFSAATVFAADNVSTAPDAVTKTLTIHKLLLSEDDLKTWD
 ORF4_14CSR MKSINKFLTMLAALLLTASSLSFSAATVFAADNVSTAPDAVTKTLTIHKLLLSEDDLKTWD
 ORF4_19AH MKSINKFLTMLAALLLTASSLSFSAATVFAADNVSTAPDAVTKTLTIHKLLLSEDDLKTWD
 ORF4_23FP MKSINKFLTMLAALLLTASSLSFSAATVFAADNVSTAPDAVTKTLTIHKLLLSEDDLKTWD
 ORF4_23FTW MKSINKFLTMLAALLLTASSLSFSAATVFAADNVSTAPDAVTKTLTIHKLLLSEDDLKTWD
 ORF4_19FTW MKSINKFLTMLAALLLTASSLSFSAATVFAAEQK-----TKTLTVHKLLMTDQELDawn
 ORF4_9VSP MKSINKFLTMLAALLLTASSLSFSAATVFAAGTT-----TTSVTVHKLLATDGDMDKIA
 ORF4_TIGR MKSINKFLTMLAALLLTASSLSFSAATVFAAGTT-----TTSVTVHKLLATDGDMDKIA
 *****:*****.*****
 *:*** ::::

ORF4_6BF TNGPK--GYDGTQ-----SSLKDLTGVA--EEIPNVYFELQKYNLTGGEKENLKDD-S
 ORF4_6BSP TNGPK--GYDGTQ-----SSLKDLTGVA--EEIPNVYFELQKYNLTGGEKENLKDD-S
 ORF4_670 TNGPK--GYDGTQ-----SSLKDLTGVA--EEIPNVYFELQKYNLTGGEKENLKDD-S
 ORF4_14CSR TNGPK--GYDGTQ-----SSLKDLTGVA--EEIPNVYFELQKYNLTGGEKENLKDD-S
 ORF4_19AH TNGPK--GYDGTQ-----SSLKDLTGVA--EEIPNVYFELQKYNLTGGEKENLKDD-S
 ORF4_23FP TNGPK--GYDGTQ-----SSLKDLTGVA--EEIPNVYFELQKYNLTGGEKENLKDD-S
 ORF4_23FTW TNGPK--GYDGTQ-----SSLKDLTGVA--EEIPNVYFELQKYNLTGGEKENLKDD-S
 ORF4_19FTW SDAITTAGYDGSQN----FEQFQKQLQGVPGVTEISGVAFELQSYTGPQGKEQENLT-A
 ORF4_9VSP NELETG-NYAGNKVGVLPANAKEIAGVMFVWNTNTNNEIIDENGQTLGVNIDPQTFKLSGA
 ORF4_TIGR NELETG-NYAGNKVGVLPANAKEIAGVMFVWNTNTNNEIIDENGQTLGVNIDPQTFKLSGA
 :. . * *: . *: * : . : : . : : : :

ORF4_6BF KWTTVHGGLTTKDGLKIETSTLKG-VYRIREDRTKTTYVGPNGQVLTGSKAVPALVTLPL
 ORF4_6BSP KWTTVHGGLTTKDGLKIETSTLKG-VYRIREDRTKTTYVGPNGQVLTGSKAVPALVTLPL
 ORF4_670 KWTTVHGGLTTKDGLKIETSTLKG-VYRIREDRTKTTYVGPNGQVLTGSKAVPALVTLPL
 ORF4_14CSR KWTTVHGGLTTKDGLKIETSTLKG-VYRIREDRTKTTYVGPNGQVLTGSKAVPALVTLPL
 ORF4_19AH KWTTVHGGLTTKDGLKIETSTLKG-VYRIREDRTKTTYVGPNGQVLTGSKAVPALVTLPL
 ORF4_23FP KWTTVHGGLTTKDGLKIETSTLKG-VYRIREDRTKTTYVGPNGQVLTGSKAVPALVTLPL
 ORF4_23FTW KWTTVHGGLTTKDGLKIETSTLKG-VYRIREDRTKTTYVGPNGQVLTGSKAVPALVTLPL
 ORF4_19FTW VWTAVNKGVTTETGVKFDTEVLQG-TYRLVEVRKESTYVGPNGKVLTMKAVPALITLPL
 ORF4_9VSP MPATAMKKLTEAEGAKFNTANLPAKYKIYEIHSLSYVGEDGATLTGSKAVPIEIELPL
 ORF4_TIGR MPATAMKKLTEAEGAKFNTANLPAKYKIYEIHSLSYVGEDGATLTGSKAVPIEIELPL
 :. : * * *: * *: * : * : * : * : * : * : * : * :

ORF4_6BF VNNNGTVIDAHVFPKNSYNKPVVDKRIADTLNYND-----QNGLSIGTKIPYVNTTI
 ORF4_6BSP VNNNGTVIDAHVFPKNSYNKPVVDKRIADTLNYND-----QNGLSIGTKIPYVNTTI
 ORF4_670 VNNNGTVIDAHVFPKNSYNKPVVDKRIADTLNYND-----QNGLSIGTKIPYVNTTI
 ORF4_14CSR VNNNGTVIDAHVFPKNSYNKPVVDKRIADTLNYND-----QNGLSIGTKIPYVNTTI
 ORF4_19AH VNNNGTVIDAHVFPKNSYNKPVVDKRIADTLNYND-----QNGLSIGTKIPYVNTTI
 ORF4_23FP VNNNGTVIDAHVFPKNSYNKPVVDKRIADTLNYND-----QNGLSIGTKIPYVNTTI
 ORF4_23FTW VNNNGTVIDAHVFPKNSYNKPVVDKRIADTLNYND-----QNGLSIGTKIPYVNTTI
 ORF4_19FTW VNNNGTVIDAHVFPKNSYEDKPTATKTFDTAAGFVDP----GEKLAIGTKVPIVNTTI
 ORF4_9VSP ND----VVDHVPKNTAKPKIDKDFKGANPDTPRVDKDPVNHQVGDVVEYEVTKI
 ORF4_TIGR ND----VVDHVPKNTAKPKIDKDFKGANPDTPRVDKDPVNHQVGDVVEYEVTKI
 : * : * : * : * : * : * : * : * : * :

ORF4_6BF PSNATFATSEFSDMTEGLTYN-EDVTITLNNVAMDQADYEVTKGNNGFNLKLTEAGLAK
 ORF4_6BSP PSNATFATSEFSDMTEGLTYN-EDVTITLNNVAMDQADYEVTKGNNGFNLKLTEAGLAK
 ORF4_670 PSNATFATSEFSDMTEGLTYN-EDVTITLNNVAMDQADYEVTKGNNGFNLKLTEAGLAK
 ORF4_14CSR PSNATFATSEFSDMTEGLTYN-EDVTITLNNVAMDQADYEVTKGNNGFNLKLTEAGLAK
 ORF4_19AH PSNATFATSEFSDMTEGLTYN-EDVTITLNNVAMDQADYEVTKGNNGFNLKLTEAGLAK
 ORF4_23FP PSNATFATSEFSDMTEGLTYN-EDVTITLNNVAMDQADYEVTKGNNGFNLKLTEAGLAK
 ORF4_23FTW PSNATFATSEFSDMTEGLTYN-EDVTITLNNVAMDQADYEVTKGNNGFNLKLTEAGLAK
 ORF4_19FTW PKNSTLATAFSDMTEGLDYN-GDVVNYNGQPLDNSHYTLEAGHNGFILKLEAGLEA
 ORF4_9VSP PALANYATANWSDRMTEGLAFNKGTVKVTVDVALEAGDYALTEVATGFDLKLTDAGLAK
 ORF4_TIGR PALANYATANWSDRMTEGLAFNKGTVKVTVDVALEAGDYALTEVATGFDLKLTDAGLAK
 * :. : * : * : * : * : * : * : * : * : * :

Figure 143B

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ORF4_6BF      INGKDADQKIQTYSATLNSLAVADIPESNDITYHYGNHQDHGNTPKPTKPN-NGQITVT
ORF4_6BSP     INGKDADQKIQTYSATLNSLAVADIPESNDITYHYGNHQDHGNTPKPTKPN-NGQITVT
ORF4_670      INGKDADQKIQTYSATLNSLAVADIPESNDITYHYGNHQDHGNTPKPTKPN-NGQITVT
ORF4_14CSR    INGKDADQKIQTYSATLNSLAVADIPESNDITYHYGNHQDHGNTPKPTKPN-NGQITVT
ORF4_19AH     INGKDADQKIQTYSATLNSLAVADIPESNDITYHYGNHQDHGNTPKPTKPN-NGQITVT
ORF4_23FP     INGKDADQKIQTYSATLNSLAVADIPESNDITYHYGNHQDHGNTPKPTKPN-NGQITVT
ORF4_23FTW    INGKDAEATITLKYTATLNALAVADVPEANDVTFHYGNPNPHGNTPKPNKPK-NGELTIT
ORF4_19FTW    VNDQNAEKTVKITYSATLNDKKAIVEVPESNDVTFNYGNPNPDHGNTPKPNKPNENGDLTLT
ORF4_9VSP     VNDQNAEKTVKITYSATLNDKKAIVEVPESNDVTFNYGNPNPDHGNTPKPNKPNENGDLTLT
ORF4_TIGR     VNDQNAEKTVKITYSATLNDKKAIVEVPESNDVTFNYGNPNPDHGNTPKPNKPNENGDLTLT
:*.::*: : :.::***** *::::*****:*****:*****:*****:*****:

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ORF4_6BF      KTWDSQPAP---EGVKATVQLVNAKTGEKVGAP-----VELSENNWTTYTWSGLDNSIEY
ORF4_6BSP     KTWDSQPAP---EGVKATVQLVNAKTGEKVGAP-----VELSENNWTTYTWSGLDNSIEY
ORF4_670      KTWDSQPAP---EGVKATVQLVNAKTGEKVGAP-----VELSENNWTTYTWSGLDNSIEY
ORF4_14CSR    KTWDSQPAP---EGVKATVQLVNAKTGEKVGAP-----VELSENNWTTYTWSGLDNSIEY
ORF4_19AH     KTWDSQPAP---EGVKATVQLVNAKTGEKVGAP-----VELSENNWTTYTWSGLDNSIEY
ORF4_23FP     KTWDSQPAP---EGVKATVQLVNAKTGEKVGAP-----VELSENNWTTYTWSGLDNSIEY
ORF4_23FTW    KTWADAKDAPI-AGVEVTFDLVNAQTGEVVKVPGHETGIVLNQTNNWTFATGLDNNTTEY
ORF4_19FTW    KTWVDATGAPIPAGAEATFDLVNAQTGKVVQTV-----TLTTDKNTVTVNGLDKNTEY
ORF4_9VSP     KTWVDATGAPIPAGAEATFDLVNAQTGKVVQTV-----TLTTDKNTVTVNGLDKNTEY
ORF4_TIGR     KTWVDATGAPIPAGAEATFDLVNAQTGKVVQTV-----TLTTDKNTVTVNGLDKNTEY
***          *:*:*:*:*:*:*:*:*:*:*:*:*:*:*:*:*:*:*:*:*:*:*:*

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ORF4_6BF      K-VEEEYNGYSAEY--TVESKGLGVKNWKDNNPAPINPEEPVKTYGKKFVKVDQKDTRL
ORF4_6BSP     K-VEEEYNGYSAEY--TVESKGLGVKNWKDNNPAPINPEEPVKTYGKKFVKVDQKDTRL
ORF4_670      K-VEEEYNGYSAEY--TVESKGLGVKNWKDNNPAPINPEEPVKTYGKKFVKVDQKDTRL
ORF4_14CSR    K-VEEEYNGYSAEY--TVESKGLGVKNWKDNNPAPINPEEPVKTYGKKFVKVDQKDTRL
ORF4_19AH     K-VEEEYNGYSAEY--TVESKGLGVKNWKDNNPAPINPEEPVKTYGKKFVKVDQKDTRL
ORF4_23FP     K-VEEEYNGYSAEY--TVESKGLGVKNWKDNNPAPINPEEPVKTYGKKFVKVDQKDTRL
ORF4_23FTW    KFVERTIKGYSADYQTITETGKIAVKNWKDENPEPINPEEPVKTYGKKFVKVDQKDERL
ORF4_19FTW    KFVERSIKGYSDYQEITTAGEIAVKNWKDENPKPLDPTEPKVVTYGKKFVKVNDKDNRL
ORF4_9VSP     KFVERSIKGYSDYQEITTAGEIAVKNWKDENPKPLDPTEPKVVTYGKKFVKVNDKDNRL
ORF4_TIGR     KFVERSIKGYSDYQEITTAGEIAVKNWKDENPKPLDPTEPKVVTYGKKFVKVNDKDNRL
* * * * * : * * * * * : * * * * * : * * * * * : * * * * * : * * * * *

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```
ORF4_6BF      ENAQFVVKKADSN-KYIAFKSTAQQAADEKAAATAKQKLDAAVAAY---TNAADKQAAQA
ORF4_6BSP     ENAQFVVKKADSN-KYIAFKSTAQQAADEKAAATAKQKLDAAVAAY---TNAADKQAAQA
ORF4_670      ENAQFVVKKADSN-KYIAFKSTAQQAADEKAAATAKQKLDAAVAAY---TNAADKQAAQA
ORF4_14CSR    ENAQFVVKKADSN-KYIAFKSTAQQAADEKAAATAKQKLDAAVAAY---TNAADKQAAQA
ORF4_19AH     ENAQFVVKKADSN-KYIAFKSTAQQAADEKAAATAKQKLDAAVAAY---TNAADKQAAQA
ORF4_23FP     ENAQFVVKKADSN-KYIAFKSTAQQAADEKAAATAKQKLDAAVAAY---TNAADKQAAQA
ORF4_23FTW    KEAQFVVKNEQG--KYLALKSAAQAVNEKAAAEEKQALDAAIAAY---TNAADKNAAQA
ORF4_19FTW    AGAEFVIANADNAGQYLARKADKVSQEEKQLVVTTTKDALDRAVAAYNALTAQQQTQQEKE
ORF4_9VSP     AGAEFVIANADNAGQYLARKADKVSQEEKQLVVTTTKDALDRAVAAYNALTAQQQTQQEKE
ORF4_TIGR     AGAEFVIANADNAGQYLARKADKVSQEEKQLVVTTTKDALDRAVAAYNALTAQQQTQQEKE
               *:::*::*:::*:::*:::*:::*:::*:::*:::*
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```

ORF4_6BF      LVDQAQQEYNVAYKEAKFGYEVAGKDE--AMVLTSNTDGGFQISGLAAGT  KRLLETKAL
ORF4_6BSP     LVDQAQQEYNVAYKEAKFGYEVAGKDE--AMVLTSNTDGGFQISGLAAGT  KRLLETKAL
ORF4_670      LVDQAQQEYNVAYKEAKFGYEVAGKDE--AMVLTSNTDGGFQISGLAAGT  KRLLETKAL
ORF4_14CSR    LVDQAQQEYNVAYKEAKFGYEVAGKDE--AMVLTSNTDGGFQISGLAAGT  KRLLETKAL
ORF4_19AH     LVDQAQQEYNVAYKEAKFGYEVAGKDE--AMVLTSNTDGGFQISGLAAGT  KRLLETKAL
ORF4_23FP     LVDQAQQEYNVAYKEAKFGYEVAGKDE--AMVLTSNTDGGFQISGLAAGT  KRLLETKAL
ORF4_23FTW    VVDAAQKTYNDNYRAARFGYEVERKED--ALVLTSNTDGGFQISGLAAGS  KRLLETKAL
ORF4_19FTW    KVDKAQAAYNAAVIAANNAFEWVADKDNENVVKLVSDAQGRFEITGLLAGT  KRLLETKAL
ORF4_9VSP     KVDKAQAAYNAAVIAANNAFEWVADKDNENVVKLVSDAQGRFEITGLLAGT  KRLLETKAL
ORF4_TIGR     KVDKAQAAYNAAVIAANNAFEWVADKDNENVVKLVSDAQGRFEITGLLAGT  KRLLETKAL

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PCT/US2005/027239

Figure 143C

ORF4_6BF	AKIDD-VEFVVGAGSWNQG--EFNYLKDVQKNDATKVVNKKITIPOTGGIGTIIFAV
ORF4_6BSP	AKIDD-VEFVVGAGSWNQG--EFNYLKDVQKNDATKVVNKKITIPOTGGIGTIIFAV
ORF4_670	AKIDD-VEFVVGAGSWNQG--EFNYLKDVQKNDATKVVNKKITIPOTGGIGTIIFAV
ORF4_14CSR	AKIDD-VEFVVGAGSWNQG--EFNYLKDVQKNDATKVVNKKITIPOTGGIGTIIFAV
ORF4_19AH	AKIDD-VEFVVGAGSWNQG--EFNYLKDVQKNDATKVVNKKITIPOTGGIGTIIFAV
ORF4_23FP	AKIDD-VEFVVGAGSWNQG--EFNYLKDVQKNDATKVVNKKITIPOTGGIGTIIFAV
ORF4_23FTW	AKLGD-VKFEVGAGSWNQG--DFNYLKDVQKNDATKVVNKKITIPOTGGIGTIIFAV
ORF4_19FTW	ALLTSRQKFEVTATSYSATGQGIETAGSGKDDATKVVNKKITIPOTGGIGTIIFAV
ORF4_9VSP	ALLTSRQKFEVTATSYSATGQGIETAGSGKDDATKVVNKKITIPOTGGIGTIIFAV
ORF4_TIGR	ALLTSRQKFEVTATSYSATGQGIETAGSGKDDATKVVNKKITIPOTGGIGTIIFAV

: : . : * * * * : . : * . * :*****

ORF4_6BF	AGAAIMGIAVYAYVKNNKDEDQLA
ORF4_6BSP	AGAAIMGIAVYAYVKNNKDEDQLA
ORF4_670	AGAAIMGIAVYAYVKNNKDEDQLA
ORF4_14CSR	AGAAIMGIAVYAYVKNNKDEDQLA
ORF4_19AH	AGAAIMGIAVYAYVKNNKDEDQLA
ORF4_23FP	AGAVIMGIAVYAYVKNNKDEDQLA
ORF4_23FTW	AGAVIMGIAVYAYVKNNKDEDQLA
ORF4_19FTW	AGAVIMGIAVYAYVKNNKDEDQLA
ORF4_9VSP	AGAVIMGIAVYAYVKNNKDEDQLA
ORF4_TIGR	AGAAIMGIAVYAYVKNNKDEDQLA

*** . *****

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Figure 144A

ORF5_6BSP -----MTMQMKQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEV
ORF5_TIGR -----MTMQMKQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEV
ORF5_6BF -----MTMQMKQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEV
ORF5_670 -----MTMQMKQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEV
ORF5_19AH -----MTMQMKQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEV
ORF5_14CSR -----MTMQMKQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEV
ORF5_19FTW -----MTMQMKQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEV
ORF5_23FTW -----MTMQMKQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEV
ORF5_9VSP MTMQMKQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEV
ORF5_23FP -----MTMQMKQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVQLQENYQEV

ORF5_6BSP VSQLP SRDGHRLQVWKLDDSYSDRRVQIVRDLHSDENKLSSEFKKTSFEMTFLENQIEV
ORF5_TIGR VSQLP SRDGHRLQVWKLDDSYSDRRVQIVRDLHSDENKLSSEFKKTSFEMTFLENQIEV
ORF5_6BF VSQLP SRDGHRLQVWKLDDSYSDRRVQIVRDLHSDENKLSSEFKKTSFEMTFLENQIEV
ORF5_670 VSQLP SRDGHRLQVWKLDDSYSDRRVQIVRDLHSDENKLSSEFKKTSFEMTFLENQIEV
ORF5_19AH VSQLP SRDGHRLQVWKLDDSYSDRRVQIVRDLHSDENKLSSEFKKTSFEMTFLENQIEV
ORF5_14CSR VSQLP SRDGHRLQVWKLDDSYSDRRVQIVRDLHSDENKLSSEFKKTSFEMTFLENQIEV
ORF5_19FTW VSQLP SRDGHRLQVWKLDDSYSDRRVQIVRDLHSDENKLSSEFKKTSFEMTFLENQIEV
ORF5_23FTW VSQLP SRDGHRLQVWKLDDSYSDRRVQIVRDLHSDENKLSSEFKKTSFEMTFLENQIEV
ORF5_9VSP VSQLP SRDGHRLQVWKLDDSYSDRRVQIVRDLHSDENKLSSEFKKTSFEMTFLENQIEV
ORF5_23FP VSQLP SRDGHRLQVWKLDDSYSDRRVQIVRDLHSDENKLSSEFKKTSFEMTFLENQIEV

ORF5_6BSP SHIPNGLYYVRSIIQTDVASYPAEFLFEMTDQTVPLVIVAKKTDMTTKVKLIKVDQDH
ORF5_TIGR SHIPNGLYYVRSIIQTDVASYPAEFLFEMTDQTVPLVIVAKKTDMTTKVKLIKVDQDH
ORF5_6BF SHIPNGLYYVRSIIQTDVASYPAEFLFEMTDQTVPLVIVAKKTDMTTKVKLIKVDQDH
ORF5_670 SHIPNGLYYVRSIIQTDVASYPAEFLFEMTDQTVPLVIVAKKTDMTTKVKLIKVDQDH
ORF5_19AH SHIPNGLYYVRSIIQTDVASYPAEFLFEMTDQTVPLVIVAKKTDMTTKVKLIKVDQDH
ORF5_14CSR SHIPNGLYYVRSIIQTDVASYPAEFLFEMTDQTVPLVIVAKKTDMTTKVKLIKVDQDH
ORF5_19FTW SHIPNGLYYVRSIIQTDVASYPAEFLFEMTDQTVPLVIVAKKTDMTTKVKLIKVDQDH
ORF5_23FTW SHIPNGLYYVRSIIQTDVASYPAEFLFEMTDQTVPLVIVAKKTDMTTKVKLIKVDQDH
ORF5_9VSP SHIPNGLYYVRSIIQTDVASYPAEFLFEMTDQTVPLVIVAKKTDMTTKVKLIKVDQDH
ORF5_23FP SHIPNGLYYVRSIIQTDVASYPAEFLFEMTDQTVPLVIVAKKTDMTTKVKLIKVDQDH

ORF5_6BSP NRLEGVGFKLVS VARDGSEKEVPLIGEYRYS SSGQVGR TLYTDKNGE I FVTNLPLGN YRF
ORF5_TIGR NRLEGVGFKLVS VARDVSEKEVPLIGEYRYS SSGQVGR TLYTDKNGE I FVTNLPLGN YRF
ORF5_6BF NRLEGVGFKLVS VARDGSEKEVPLIGEYRYS SSGQVGR TLYTDKNGE I FVTNLPLGN YRF
ORF5_670 NRLEGVGFKLVS VARDGSEKEVPLIGEYRYS SSGQVGR TLYTDKNGE I FVTNLPLGN YRF
ORF5_19AH NRLEGVGFKLVS VARDGSEKEVPLIGEYRYS SSGQVGR TLYTDKNGE I FVTNLPLGN YRF
ORF5_14CSR NRLEGVGFKLVS VARDGSEKEVPLIGEYRYS SSGQVGR TLYTDKNGE I FVTNLPLGN YRF
ORF5_19FTW NRLEGVGFKLVS VARDGSEKEVPLIGEYRYS SSGQVGR TLYTDKNGE I FVTNLPLGN YRF
ORF5_23FTW NRLEGVGFKLVS VARDGSEKEVPLIGEYRYS SSGQVGR TLYTDKNGE I FVTNLPLGN YRF
ORF5_9VSP NRLEGVGFKLVS VARDGSEKEVPLIGEYRYS SSGQVGR TLYTDKNGE I FVTNLPLGN YRF
ORF5_23FP NRLEGVGFKLVS VARDGSEKEVPLIGEYRYS SSGQVGR TLYTDKNGE I FVTNLPLGN YRF

ORF5_6BSP KEVEPLAGYAVTTLD TDVQLVDHQLVTITVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKV
ORF5_TIGR KEVEPLAGYAVTTLD TDVQLVDHQLVTITVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKV
ORF5_6BF KEVEPLAGYAVTTLD TDVQLVDHQLVTITVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKV
ORF5_670 KEVEPLAGYAVTTLD TDVQLVDHQLVTITVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKV
ORF5_19AH KEVEPLAGYAVTTLD TDVQLVDHQLVTITVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKV
ORF5_14CSR KEVEPLAGYAVTTLD TDVQLVDHQLVTITVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKV
ORF5_19FTW KEVEPLAGYAVTTLD TDVQLVDHQLVTITVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKV
ORF5_23FTW KEVEPLAGYAVTTLD TDVQLVDHQLVTITVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKV
ORF5_9VSP KEVEPLAGYAVTTLD TDVQLVDHQLVTITVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKV
ORF5_23FP KEVEPLAGYAVTTLD TDVQLVDHQLVTITVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKV

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Figure 144B

```
ORF5_6BSP      MKEESGHYTPVLQNGKEVVVTSGKDGRFRVEGLE YGT  LWELOAPTCGVQLTSPVSFTI
ORF5_TIGR      MKEESGHYTPVLQNGKEVVVTSGKDGRFRVEGLE YGT  LWELOAPTCGVQLTSPVSFTI
ORF5_6BF       MKEESGHYTPVLQNGKEVVVTSGKDGRFRVEGLE YGT  LWELOAPTCGVQLTSPVSFTI
ORF5_670       MKEESGHYTPVLQNGKEVVVTSGKDGRFRVEGLE YGT  LWELOAPTCGVQLTSPVSFTI
ORF5_19AH      MKEESGHYTPVLQNGKEVVVTSGKDGRFRVEGLE YGT  LWELOAPTCGVQLTSPVSFTI
ORF5_14CSR     MKEESGHYTPVLQNGKEVVVTSGKDGRFRVEGLE YGT  LWELOAPTCGVQLTSPVSFTI
ORF5_19FTW     MKEENGHYTPVLQNGKEVVVASGKDGRFRVEGLE YGT  LWELOAPTCGVQLTSPVSFTI
ORF5_23FTW     MKEENGHYTPVLQNGKEVVVASGKDGRFRVEGLE YGT  LWELOAPTCGVQLTSPVSFTI
ORF5_9VSP      MKEENGHYTPVLQNGKEVVVASGKDGRFRVEGLE YGT  LWELOAPTCGVQLTSPVSFTI
ORF5_23FP      MKEENGHYTPVLQNGKEVVVASGKDGRFRVEGLE YGT  LWELOAPTCGVQLTSPVSFTI
*****
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```
ORF5_6BSP      GKDTRKELVTVVKNNKRPRIDV P D T G E E T L Y I L M L V A I L L F G S G Y Y L T K K P N N
ORF5_TIGR      GKDTRKELVTVVKNNKRPRIDV P D T G E E T L Y I L M L V A I L L F G S G Y Y L T K K P N N
ORF5_6BF       GKDTRKELVTVVKNNKRPRIDV P D T G E E T L Y I L M L V A I L L F G S G Y Y L T K K P N N
ORF5_670       GKDTRKELVTVVKNNKRPRIDV P D T G E E T L Y I L M L V A I L L F G S G Y Y L T K K P N N
ORF5_19AH      GKDTRKELVTVVKNNKRPRIDV P D T G E E T L Y I L M L V A I L L F G S G Y Y L T K K P N N
ORF5_14CSR     GKDTRKELVTVVKNNKRPRIDV P D T G E E T L Y I L M L V A I L L F G S G Y Y L T K K P N N
ORF5_19FTW     GKDTRKELVTVVKNNKRPRIDV P D T G E E T L Y I L M L V A I L L F G S G Y Y L T K K T N N
ORF5_23FTW     GKDTRKELVTVVKNNKRPRIDV P D T G E E T L Y I L M L V A I L L F G S G Y Y L T K K T N N
ORF5_9VSP      GKDTRKELVTVVKNNKRPRIDV P D T G E E T L Y I L M L V A I L L F G S G Y Y L T K K T N N
ORF5_23FP      GKDTRKELVTVVKNNKRPRIDV P D T G E E T L Y I L M L V A I L L F G S G Y Y L T K K T N N
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Figure 145A

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ORF6_23FTW      MLIKMKVTKKQKRNNLLLGTVFFIGMAVMAYPLVSRLYYRVESNQIADFDKEKATLDEA
ORF6_TIGR       MLIKMKVTKKQKRNNLLLGTVFFIGMAVMAYPLVSRLYYRVESNQIADFDKEKATLDEA
ORF6_6BSP       MLIKMKVTKKQKRNNLLLGTVFFIGMAVMAYPLVSRLYYRVESNQIADFDKEKATLDEA
ORF6_6BF        MLIKMKVTKKQKRNNLLLGTVFFIGMAVMAYPLVSRLYYRVESNQIADFDKEKATLDEA
ORF6_670        MLIKMKVTKKQKRNNLLLGTVFFIGMAVMAYPLVSRLYYRVESNQIADFDKEKATLDEA
ORF6_19AH       MLIKMKVTKKQKRNNLLLGTVFFIGMAVMAYPLVSRLYYRVESNQIADFDKEKATLDEA
ORF6_14CSR      MLIKMKVTKKQKRNNLLLGTVFFIGMAVMAYPLVSRLYYRVESNQIADFDKEKATLDEA
ORF6_23FP       MLIKMAKTKKQKRNNLLLGTVFFIGIAVMAYPLVSRLYYRVESNQIADFDKEKATLDEA
ORF6_9VSP       MLIKMAKTKKQKRNNLLLGTVFFIGIAVMAYPLVSRLYYRVESNQIADFDKEKATLDEA
ORF6_19FTW      MLIKMAKTKKQKRNNLLLGTVFFIGMAVMAYPLVSRLYYRVESNQIADFDKEKATLDEA
*****

ORF6_23FTW      DIDERMKLAQAFNDSLNNVSGDPWSEEMKKKGRAEYARMLEIHERMGHVEIPVIDVDLP
ORF6_TIGR       DIDERMKLAQAFNDSLNNVSGDPWSEEMKKKGRAEYARMLEIHERMGHVEIPVIDVDLP
ORF6_6BSP       DIDERMKLAQAFNDSLNNVSGDPWSEEMKKKGRAEYARMLEIHERMGHVEIPVIDVDLP
ORF6_6BF        DIDERMKLAQAFNDSLNNVSGDPWSEEMKKKGRAEYARMLEIHERMGHVEIPVIDVDLP
ORF6_670        DIDERMKLAQAFNDSLNNVSGDPWSEEMKKKGRAEYARMLEIHERMGHVEIPVIDVDLP
ORF6_19AH       DIDERMKLAQAFNDSLNNVSGDPWSEEMKKKGRAEYARMLEIHERMGHVEIPVIDVDLP
ORF6_14CSR      DIDERMKLAQAFNDSLNNVSGDPWSEEMKKKGRAEYARMLEIHERMGHVEIPVIDVDLP
ORF6_23FP       DIDERMKLAQAFNDSLNNVSGDPWSEEMKKKGRAEYARMLEIHERMGHVEIPVIDVDLP
ORF6_9VSP       DIDERMKLAQAFNDSLNNVSGDPWSEEMKKKGRAEYARMLEIHERMGHVEIPVIDVDLP
ORF6_19FTW      DIDERMKLAQAFNDSLNNVSGDPWSEEMKKKGRAEYARMLEIHERMGHVEIPVIDVDLP
*****

ORF6_23FTW      VYAGTAEVLQQGAGHLEGTSLPIGGNSTHAVITAHTGLPTAKMFTDLTKLVGDKFYVH
ORF6_TIGR       VYAGTAEVLQQGAGHLEGTSLPIGGNSTHAVITAHTGLPTAKMFTDLTKLVGDKFYVH
ORF6_6BSP       VYAGTAEVLQQGAGHLEGTSLPIGGNSTHAVITAHTGLPTAKMFTDLTKLVGDKFYVH
ORF6_6BF        VYAGTAEVLQQGAGHLEGTSLPIGGNSTHAVITAHTGLPTAKMFTDLTKLVGDKFYVH
ORF6_670        VYAGTAEVLQQGAGHLEGTSLPIGGNSTHAVITAHTGLPTAKMFTDLTKLVGDKFYVH
ORF6_19AH       VYAGTAEVLQQGAGHLEGTSLPIGGNSTHAVITAHTGLPTAKMFTDLTKLVGDKFYVH
ORF6_14CSR      VYAGTAEVLQQGAGHLEGTSLPIGGNSTHAVITAHTGLPTAKMFTDLTKLVGDKFYVH
ORF6_23FP       VYAGTAEVLQQGAGHLEGTSLPIGGNSTHAVITAHTGLPTAKMFTDLTKLVGDKFYVH
ORF6_9VSP       VYAGTAEVLQQGAGHLEGTSLPIGGNSTHAVITAHTGLPTAKMFTDLTKLVGDKFYVH
ORF6_19FTW      VYAGTAEVLQQGAGHLEGTSLPIGGNSTHAVITAHTGLPTAKMFTDLTKLVGDKFYVH
*****

ORF6_23FTW      NIKEVMAYQVDQVKVIEPTNFDDLLIVPGHDYVTLTCTPYMINTHRLVGRHRIPIYVAE
ORF6_TIGR       NIKEVMAYQVDQVKVIEPTNFDDLLIVPGHDYVTLTCTPYMINTHRLVGRHRIPIYVAE
ORF6_6BSP       NIKEVMAYQVDQVKVIEPTNFDDLLIVPGHDYVTLTCTPYMINTHRLVGRHRIPIYVAE
ORF6_6BF        NIKEVMAYQVDQVKVIEPTNFDDLLIVPGHDYVTLTCTPYMINTHRLVGRHRIPIYVAE
ORF6_670        NIKEVMAYQVDQVKVIEPTNFDDLLIVPGHDYVTLTCTPYMINTHRLVGRHRIPIYVAE
ORF6_19AH       NIKEVMAYQVDQVKVIEPTNFDDLLIVPGHDYVTLTCTPYMINTHRLVGRHRIPIYVAE
ORF6_14CSR      NIKEVMAYQVDQVKVIEPTNFDDLLIVPGHDYVTLTCTPYMINTHRLVGRHRIPIYVAE
ORF6_23FP       NIKEVMAYQVDQVKVIEPTNFDDLLIVPGHDYVTLTCTPYMINTHRLVGRHRIPIYVAE
ORF6_9VSP       NIKEVMAYQVDQVKVIEPTNFDDLLIVPGHDYVTLTCTPYMINTHRLVGRHRIPIYVAE
ORF6_19FTW      NIKEVMAYQVDQVKVIEPTNFDDLLIVPGHDYVTLTCTPYMINTHRLVGRHRIPIYVAE
*****

ORF6_23FTW      VEEEFIAANKLSHLYRYLFYVAVGLIVILLWIIRRLRKKKKQPEKALKALKAAARKEVKVE
ORF6_TIGR       VEEEFIAANKLSHLYRYLFYVAVGLIVILLWIIRRLRKKKKQPEKALKALKAAARKEVKVE
ORF6_6BSP       VEEEFIAANKLSHLYRYLFYVAVGLIVILLWIIRRLRKKKKQPEKALKALKAAARKEVKVE
ORF6_6BF        VEEEFIAANKLSHLYRYLFYVAVGLIVILLWIIRRLRKKKKQPEKALKALKAAARKEVKVE
ORF6_670        VEEEFIAANKLSHLYRYLFYVAVGLIVILLWIIRRLRKKKKQPEKALKALKAAARKEVKVE
ORF6_19AH       VEEEFIAANKLSHLYRYLFYVAVGLIVILLWIIRRLRKKKKQPEKALKALKAAARKEVKVE
ORF6_14CSR      VEEEFIAANKLSHLYRYLFYVAVGLIVILLWIIRRLRKKKKQPEKALKALKAAARKEVKVE
ORF6_23FP       VEEEFIAANKLSHLYRYLFYVAVGLIVILLWIIRRLRKKKKQSERALKALKEATKEVKVE
ORF6_9VSP       VEEEFIAANKLSHLYRYLFYVAVGLIVILLWIIRRLRKKKKQSERALKALKEATKEVKVE
ORF6_19FTW      VEEEFIAANKLSHLYRYLFYVAVGLIVILLWIIRRLRKKKKQSERALKALKEATKEVKVE
*****
```

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Figure 145B

ORF6_23FTW	DGQQ
ORF6_TIGR	DGQQ
ORF6_6BSP	DGQQ
ORF6_6BF	DGQQ
ORF6_670	DGQQ
ORF6_19AH	DGQQ
ORF6_14CSR	DGQQ
ORF6_23FP	DE--
ORF6_9VSP	DE--
ORF6_19FTW	DE--

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[illegible]

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ORF8_14CSR      MSKAKLQKLLGYLLMLVALVIPVYCFGQMVLQSLGQVKGHEIFSESVTADSYQEQLQRS
ORF8_19AH      MSKAKLQKLLGYLLMLVALVIPVYCFGQMVLQSLGQVKGHEIFSESVTADSYQEQLQRS
ORF8_23FTW     MSKAKLQKLLGYLLMLVALVIPVYCFGQMVLQSLGQVKGHEIFSESVTADSYQEQLQRS
ORF8_670       MSKAKLQKLLGYLLMLVALVIPVYCFGQMVLQSLGQVKGHEIFSESVTADSYQEQLQRS
ORF8_6BF       MSKAKLQKLLGYLLMLVALVIPVYCFGQMVLQSLGQVKGHEIFSESVTADSYQEQLQRS
ORF8_6BSP      MSKAKLQKLLGYLLMLVALVIPVYCFGQMVLQSLGQVKGHEIFSESVTADSYQEQLQRS
ORF8_19FTW     MSRTKLRALLGYLLMLVACLIPIYCFGQMVLQSLGQVKGHATFVKSMTEMYQEQQNHS
ORF8_23FP      MSRTKLRALLGYLLMLVACLIPIYCFGQMVLQSLGQVKGHATFVKSMTEMYQEQQNHS
ORF8_9VSP      MSRTKLRALLGYLLMLVACLIPIYCFGQMVLQSLGQVKGHATFVKSMTEMYQEQQNHS
ORF8_TIGR      MSRTKLRALLGYLLMLVACLIPIYCFGQMVLQSLGQVKGHATFVKSMTEMYQEQQNHS
                **:.*: ***** :*:*****
ORF8_14CSR      DYNQRLDSQNRIVDPFLAEGYEVNYQVSDPPDAVYGYLSIPSLIMEIPVYLGDYHHLAM
ORF8_19AH      DYNQRLDSQNRIVDPFLAEGYEVNYQVSDPPDAVYGYLSIPSLIMEIPVYLGDYHHLAM
ORF8_23FTW     DYNQRLDSQNRIVDPFLAEGYEVNYQVSDPPDAVYGYLSIPSLIMEIPVYLGDYHHLAM
ORF8_670       DYNQRLDSQNRIVDPFLAEGYEVNYQVSDPPDAVYGYLSIPSLIMEIPVYLGDYHHLAM
ORF8_6BF       DYNQRLDSQNRIVDPFLAEGYEVNYQVSDPPDAVYGYLSIPSLIMEIPVYLGDYHHLAM
ORF8_6BSP      DYNQRLDSQNRIVDPFLAEGYEVNYQVSDPPDAVYGYLSIPSLIMEIPVYLGDYHHLAM
ORF8_19FTW     AYNQRLASQNRIVDPFLAEGYEVNYQVSDPPDAVYGYLSIPSLIMEIPVYLGDYHHLGM
ORF8_23FP      AYNQRLASQNRIVDPFLAEGYEVNYQVSDPPDAVYGYLSIPSLIMEIPVYLGDYHHLGM
ORF8_9VSP      AYNQRLASQNRIVDPFLAEGYEVNYQVSDPPDAVYGYLSIPSLIMEIPVYLGDYHHLGM
ORF8_TIGR      AYNQRLASQNRIVDPFLAEGYEVNYQVSDPPDAVYGYLSIPSLIMEIPVYLGDYHHLGM
                *****
ORF8_14CSR      GLAHVDGTPLPVEGKGIRSVIAGHRAEPSHVFFRHLDDLQKVGDALYDNGQEIVEYQMMMD
ORF8_19AH      GLAHVDGTPLPVEGKGIRSVIAGHRAEPSHVFFRHLDDLQKVGDALYDNGQEIVEYQMMMD
ORF8_23FTW     GLAHVDGTPLPVEGKGIRSVIAGHRAEPSHVFFRHLDDLQKVGDALYDNGQEIVEYQMMMD
ORF8_670       GLAHVDGTPLPVEGKGIRSVIAGHRAEPSHVFFRHLDDLQKVGDALYDNGQEIVEYQMMMD
ORF8_6BF       GLAHVDGTPLPVEGKGIRSVIAGHRAEPSHVFFRHLDDLQKVGDALYDNGQEIVEYQMMMD
ORF8_6BSP      GLAHVDGTPLPVEGKGIRSVIAGHRAEPSHVFFRHLDDLQKVGDALYDNGQEIVEYQMMMD
ORF8_19FTW     GLAHVDGTPLPDGTGIRSVIAGHRAEPSHVFFRHLDDLQKVGDALYDNGQEIVEYQMMMD
ORF8_23FP      GLAHVDGTPLPDGTGIRSVIAGHRAEPSHVFFRHLDDLQKVGDALYDNGQEIVEYQMMMD
ORF8_9VSP      GLAHVDGTPLPDGTGIRSVIAGHRAEPSHVFFRHLDDLQKVGDALYDNGQEIVEYQMMMD
ORF8_TIGR      GLAHVDGTPLPDGTGIRSVIAGHRAEPSHVFFRHLDDLQKVGDALYDNGQEIVEYQMMMD
                *****:.*.*****
ORF8_14CSR      TEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAABARVA
ORF8_19AH      TEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAABARVA
ORF8_23FTW     TEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAABARVA
ORF8_670       TEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAABARVA
ORF8_6BF       TEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAABARVA
ORF8_6BSP      TEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAABARVA
ORF8_19FTW     TEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAABARVA
ORF8_23FP      TEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAABARVA
ORF8_9VSP      TEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAABARVA
ORF8_TIGR      TEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAABARVA
                *****
ORF8_14CSR      FTKEGQSVSRVATSQWLYRGLVLAFLGILFVLWKLARLLRGK
ORF8_19AH      FTKEGQSVSRVATSQWLYRGLVLAFLGILFVLWKLARLLRGK
ORF8_23FTW     FTKEGQSVSRVATSQWLYRGLVLAFLGILFVLWKLARLLRGK
ORF8_670       FTKEGQSVSRVATSQWLYRGLVLAFLGILFVLWKLARLLRGK
ORF8_6BF       FTKEGQSVSRVATSQWLYRGLVLAFLGILFVLWKLARLLRGK
ORF8_6BSP      FTKEGQSVSRVATSQWLYRGLVLAFLGILFVLWKLARLLRGK
ORF8_19FTW     FTKEGQSVSRVATSQWLYRGLVLAFLGILFVLWKLARLLRGK
ORF8_23FP      FTKEGQSVSRVATSQWLYRGLVLAFLGILFVLWKLARLLRGK
ORF8_9VSP      FTKEGQSVSRVATSQWLYRGLVLAFLGILFVLWKLARLLRGK
ORF8_TIGR      FTKEGQSVSRVATSQWLYRGLVLAFLGILFVLWKLARLLRGK
                *****

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RrgA, LPXTG

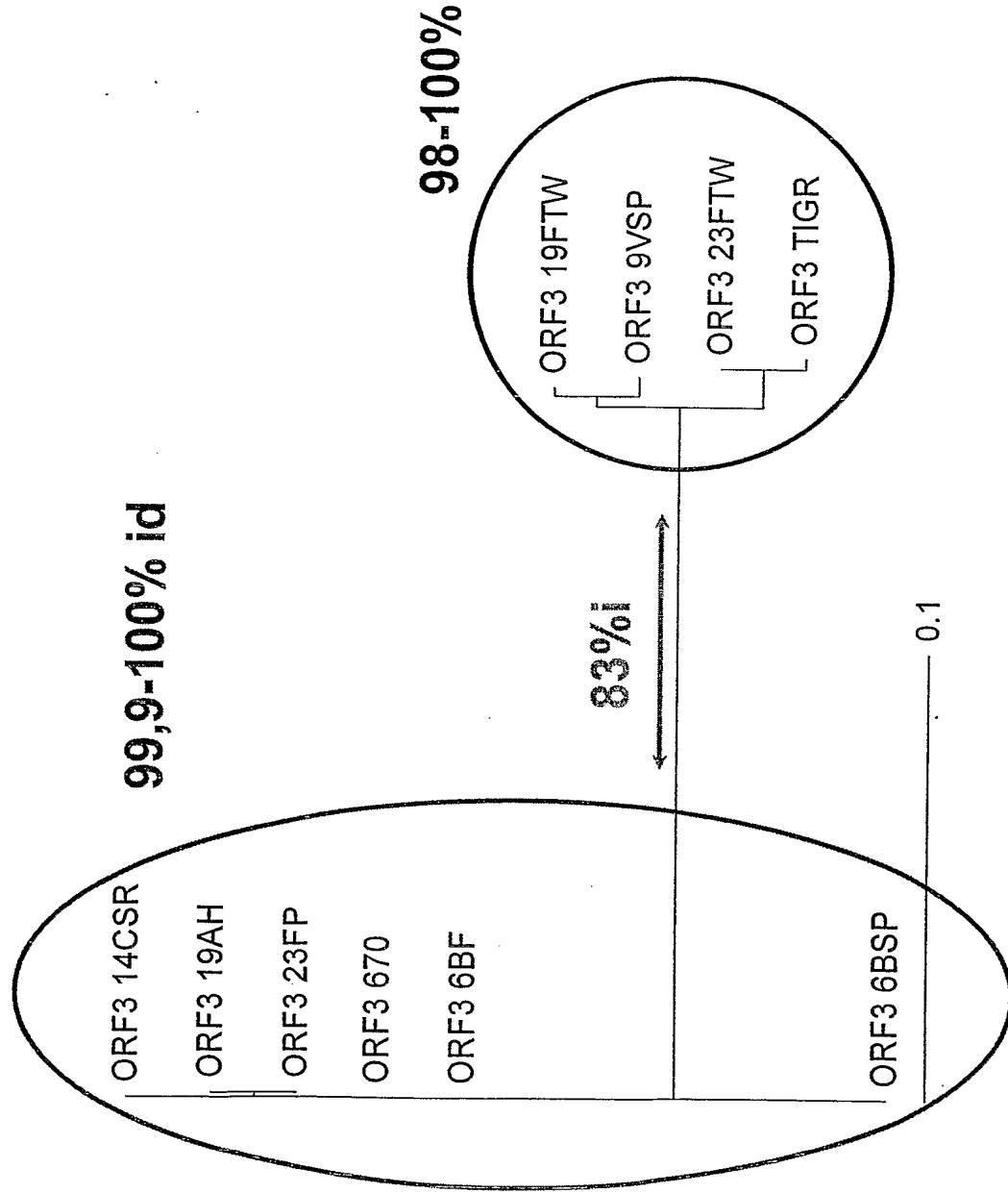


Figure 148

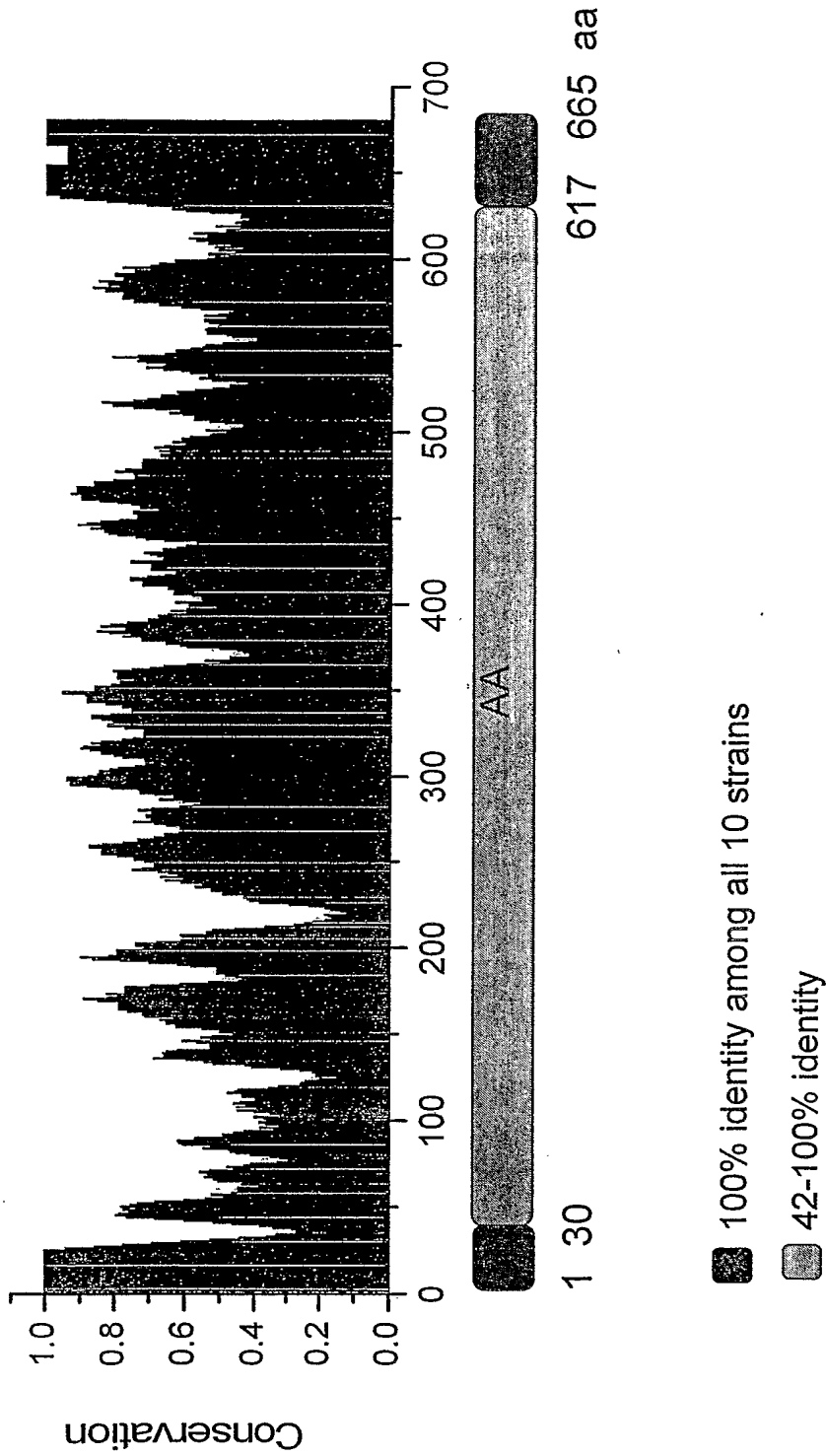


Figure 149

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A

MLNRETHMKVKRKFQKAVAGLCCISQLTAFSSIVALA*ETPETS¹PAIGKVVIKETGEGGALLGDAVFELKN
 NTDGTTVSQRTEAQTGEAIFSNIKPGTYTLTEAQQPVGYKPKSTKQWTVVEKNGRTTVQGEQVENREE
 ALSDQYPQGTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPYERVPEGTLSKRIYQVNNLDDNQYGIEL
 TVSGKTVYEQDKSVPLDVVILLDNSMSNIRNKNARRAERAGEATRS²LIDKITSSEN³RVALV⁴TYAS
 TIFDGTFTVEKGVADKNGKRLNDSLFWNYDQTSFTTNTKDYSLKLTNDKNDIVELKNKVPTAE⁵DHD
 GNRLMYQFGATFTQKALMKADEILTQQAQNSQKVIFHITDGVPTMSYPINFN⁶HATFAPS⁷YQNQLNA
 FFSKSPNKDGILLSDFITQATSGEHTTVRGDQSYQMFTDKTVYEKGAPAAFPVKPEKYSEMKAAGYAVI
 GDPINGGYIWLNWRESILAYPENSNTAKITNHGDPTRWYYNGNIAPDGYDVFTVGIGINGDPGTDEATA
 TSFMQSISSKPEN⁸YTNVTDTTKILEQLNRYFHTIVTEKKS⁹IENTITDPMGELIDLQLGTDGRFDPADYTL
 TANDGSRLENGQAVGGPQNDGGLLKNAKVL¹⁰YDTTEKRIRVTGLYLGTDEKVTLTYNVRLNDEFVSNKFYD
 TNGRTTLHPKEVEQNTVRDFPKIRDVRKYPEITISKEK¹¹LGDI¹²EIKV¹³NKN¹⁴DKK¹⁵PLRGA¹⁶VFS¹⁷LQK¹⁸QHPDYP
 DIYGAI¹⁹DQNGTYQNVRTGEDGKLTFKNLSDGKYRLFENSEPAGYKPVQNKPIVAFQIVNGEVRDVT²⁰SIVPQ
 DIPAGYEFTNDKH²¹YITNEP²²PPKREYPR²³TGGIGMLPFY²⁴LIGCMM²⁵MMGGV²⁶LLYTRKHP

B

5' cgggatcc-gaa-acg-cct-gaa-acc-agt 5' 24mer, 54 %G+C, Tm 62

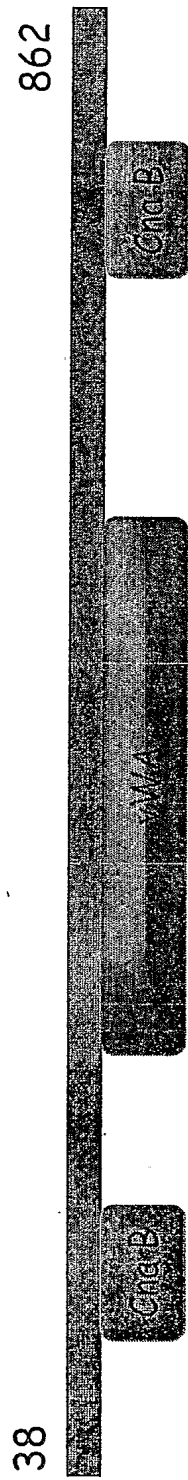
*Bam*HI

3' ccgctcgag-aat-agg-ttc-att-ggt 3' 27mer, 52 %G+C, Tm 61.6

*Xho*I

Figure 150

A.



B.

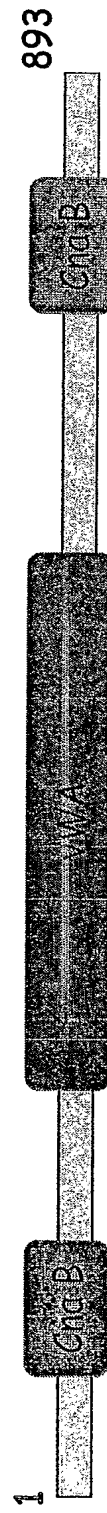
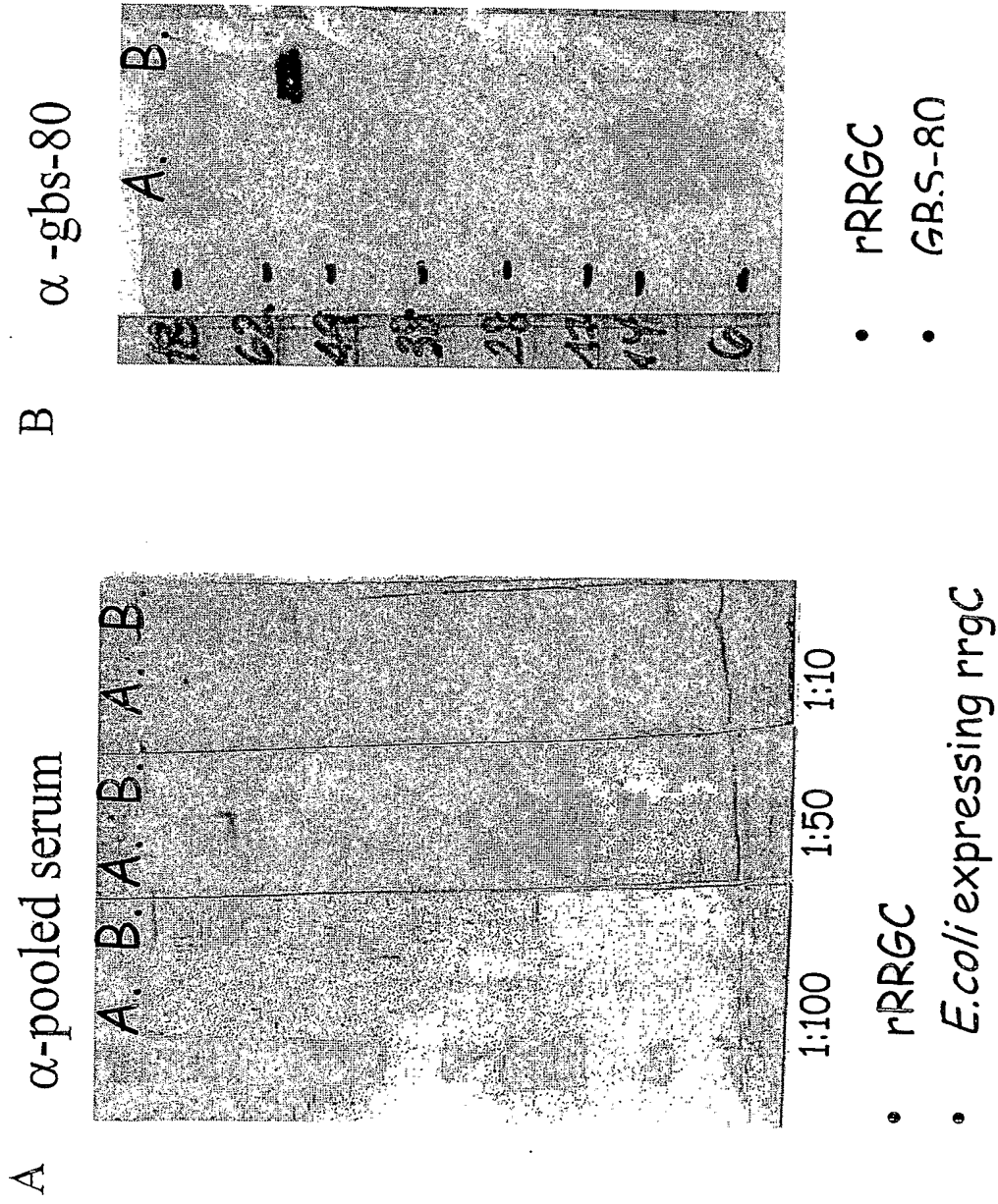


Figure 151

Figure 152



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A

MKSINKFLTMLAALLLTASSLS* AATVFAAGTTTTSVTVHKLLATDGMMDKIANELETGNYAGNKVGVLP
 NAKELAGVMFVWNTNNEIIDENGQTLGVNIDPQTFKLSGAMPATAMKKL TEAEGAKFNTANLPAAKYKIY
 EHSLS TVVGEDGATLTGSKAVPIEIELPLNDVVDVHVYPKNTAKPKIDKDFKGKANPDTPRVDKDTPVNHQV
 GDVVEYEIVTKIPALANYATANWSDRMTEGLAFNKGTVKVTVDDVVALEAGDYALTEVATGFDLKLTDAGLAK
 VNDQNAEKTIVKITYSATLNDKAIIVEVPESNDVTFNNGNPDHGNTPKNKPNENGDLTLTKTWVVDATGAPIP
 AGAEATFDLVNAQTGKVVQTVTLTTDKNTVTVNGLDKNTYKFVRSIKGYSADYQEITTA GEIAVKNNWKD
 ENPKPLDPTEPKVVITYGKKFVKVNDKDNRLAGAEFVIANADNAGQYLARKADKVSQEEKQLVVTTKDALDRAV
 AAYNALTAQQQTQQEKEKVDKAQAAYNAAVLAANNAFEWVADKDNEENVVKLVSDAQGRFEITGLLAGTY
 YLEETKQPAGYALLTSRQKFEVTATSYSATGQGIEYTAGSGKDDATKVVNKKITIPQTGGIGTIFAVAGAAI
 MGLAVYAYVKNKDEDEQLA

B

5' cgggatcc-gct-gca-aca-gtt-ttt 3' 23mer, 52.2% G+C, Tm 60.6
*Bam*HI
 5' ccgctcgcgag-agt-gat-ttt-ttt-gtt-gac 3' 26mer, 44.4% G+C, Tm 61.7
*Xho*I

Figure 153

PCT/US2005/027239 352/487

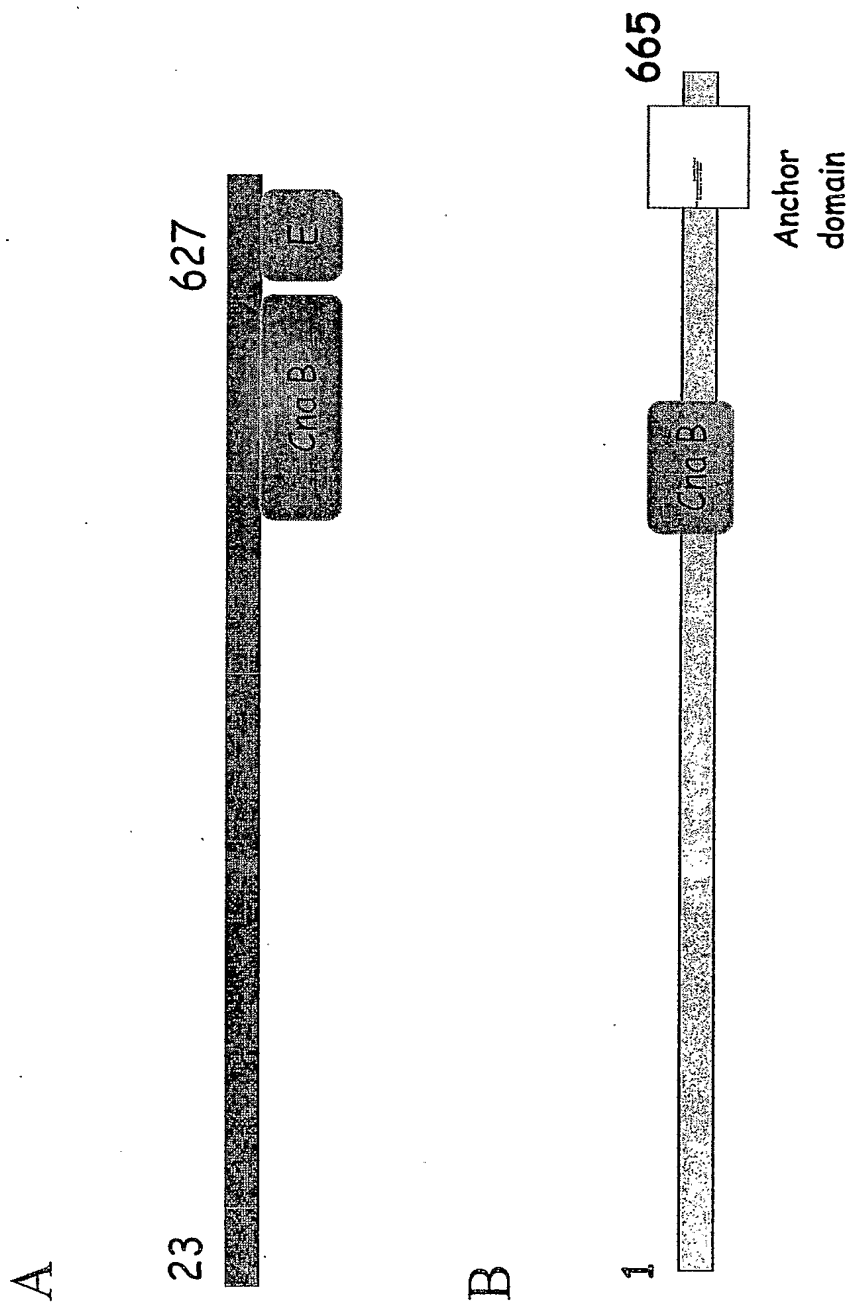


Figure 154

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60 kDa



Figure 155

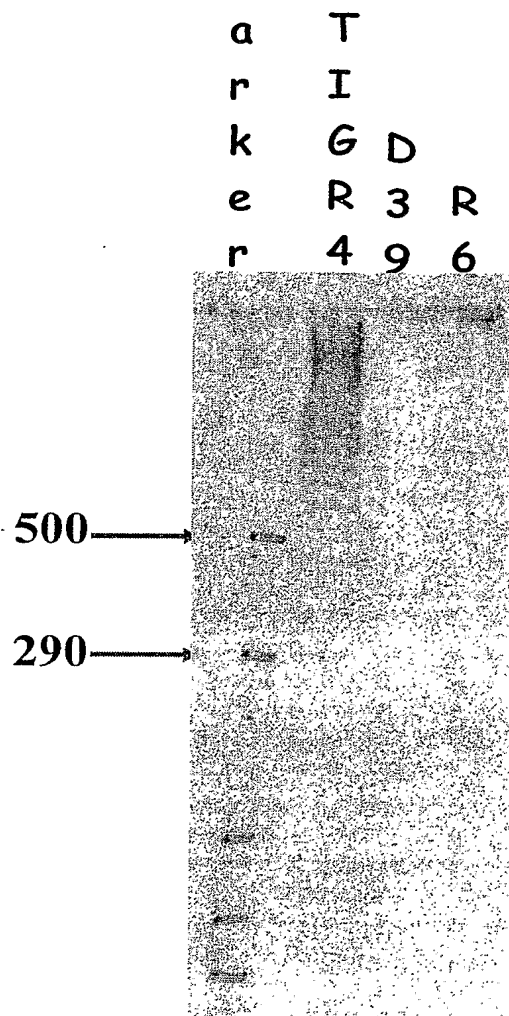


Figure 156

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A

MISRIFFV~~MALCFSLVWGA~~*H~~AVO~~Q~~AE~~D~~H~~T~~L~~V~~L~~Q~~EN~~Y~~Q~~E~~V~~V~~S~~Q~~L~~P~~S~~R~~D~~G~~H~~R~~L~~Q~~V~~W~~K~~L~~D~~D~~S~~S~~S~~
 Y~~D~~D~~R~~V~~Q~~I~~V~~R~~D~~L~~H~~S~~W~~D~~E~~N~~K~~L~~S~~S~~F~~K~~K~~T~~S~~F~~E~~M~~T~~F~~L~~E~~N~~Q~~I~~E~~V~~S~~H~~I~~P~~N~~G~~L~~Y~~Y~~V~~R~~S~~I~~Q~~T~~D~~A~~V~~S~~Y~~P~~A~~E~~F~~L~~F~~
 E~~M~~T~~D~~Q~~T~~V~~E~~P~~L~~V~~I~~V~~A~~K~~K~~T~~D~~T~~M~~T~~T~~K~~V~~K~~L~~I~~K~~V~~D~~Q~~D~~H~~N~~R~~L~~E~~G~~V~~G~~F~~K~~L~~V~~S~~V~~A~~R~~D~~V~~S~~E~~K~~E~~V~~P~~L~~I~~G~~E~~Y~~R~~Y~~S~~S~~S~~
 G~~Q~~V~~G~~R~~T~~L~~Y~~T~~D~~K~~N~~G~~E~~I~~F~~V~~T~~N~~L~~P~~L~~G~~N~~Y~~R~~F~~K~~E~~V~~E~~P~~L~~A~~G~~Y~~A~~V~~T~~T~~L~~D~~T~~D~~V~~Q~~L~~V~~D~~H~~Q~~L~~V~~T~~T~~I~~V~~V~~N~~Q~~K~~L~~P~~R~~G~~N~~
 V~~D~~F~~M~~K~~V~~D~~G~~R~~I~~N~~T~~S~~L~~Q~~G~~A~~M~~F~~K~~V~~M~~K~~E~~E~~S~~G~~H~~Y~~T~~P~~V~~L~~Q~~N~~G~~K~~E~~V~~V~~T~~S~~G~~K~~D~~G~~R~~F~~R~~V~~E~~G~~L~~E~~Y~~G~~T~~Y~~Y~~L~~W~~E~~L~~Q~~
 A~~P~~T~~G~~Y~~V~~Q~~L~~T~~S~~P~~V~~S~~F~~T~~I~~G~~K~~D~~T~~R~~K~~E~~L~~V~~T~~V~~V~~K~~N~~N~~K~~R~~P~~R~~D~~V~~P~~D~~T~~G~~E~~E~~T~~L~~Y~~I~~L~~M~~L~~V~~A~~I~~L~~L~~F~~G~~S~~G~~Y~~Y~~L~~T~~K~~K~~P~~
 NN

B

5' cgggatcc-cat-gtc-caa-gcg-caa-gaa 21mer, 61% G+C, Tm 60.8

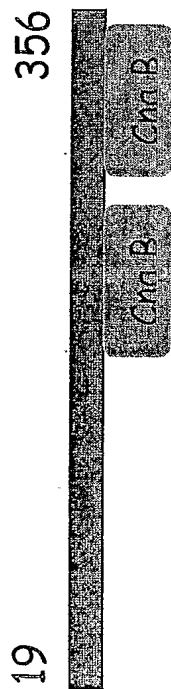
*Bam*HI

5' ccgctcgag-ctt-gtt-att-ttt-aac-cac 27mer, 44% G+C, Tm 58.4

*Xho*I

Figure 157

A



B



Figure 158

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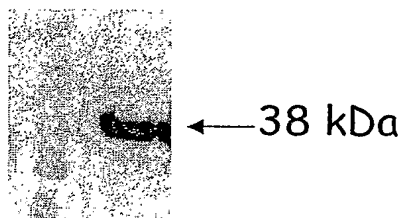


Figure 159

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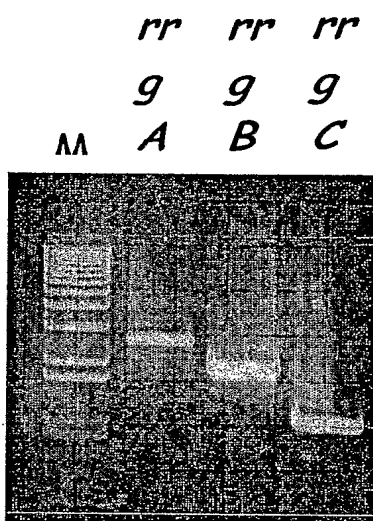
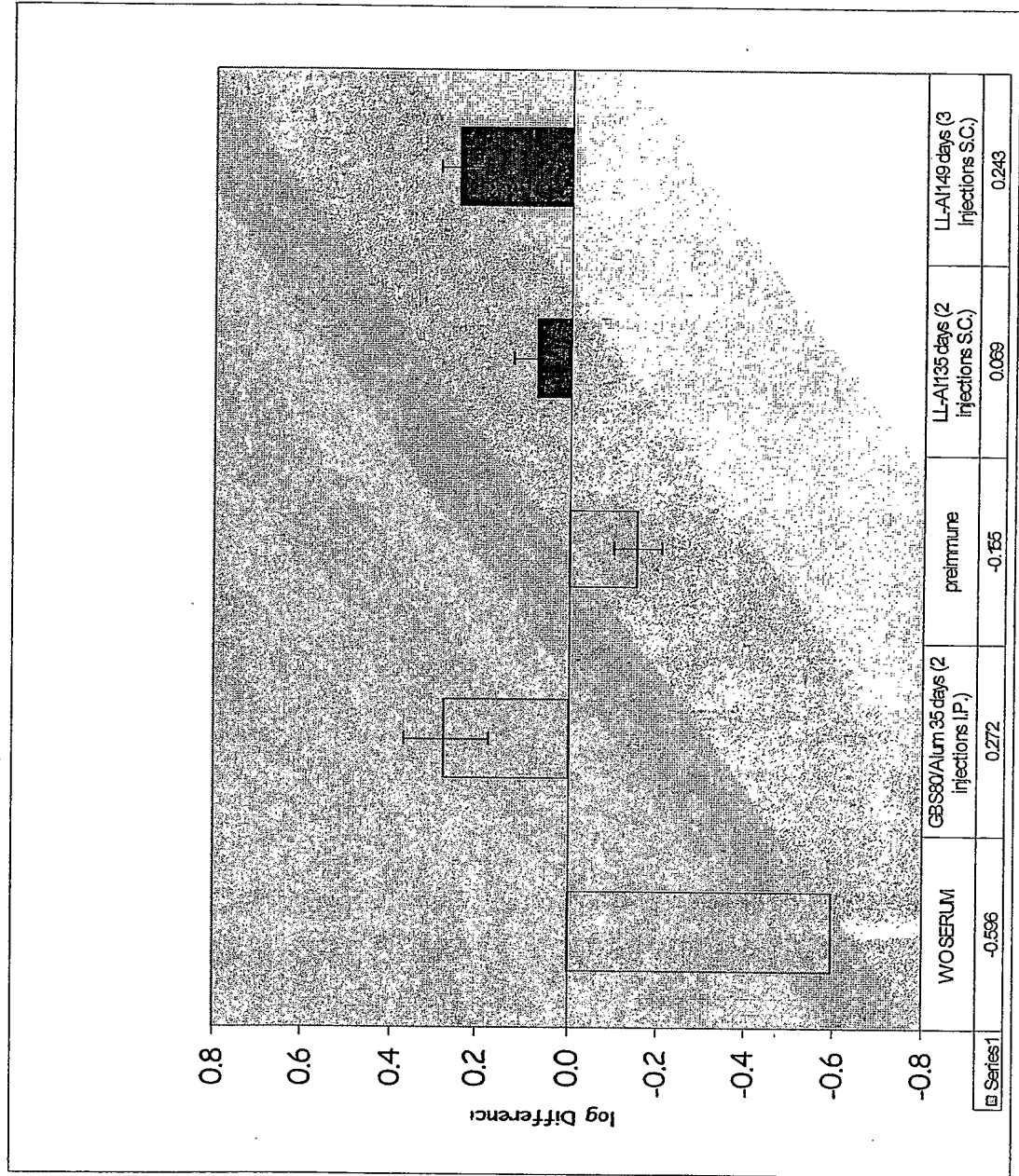


Figure 160

Figure 161



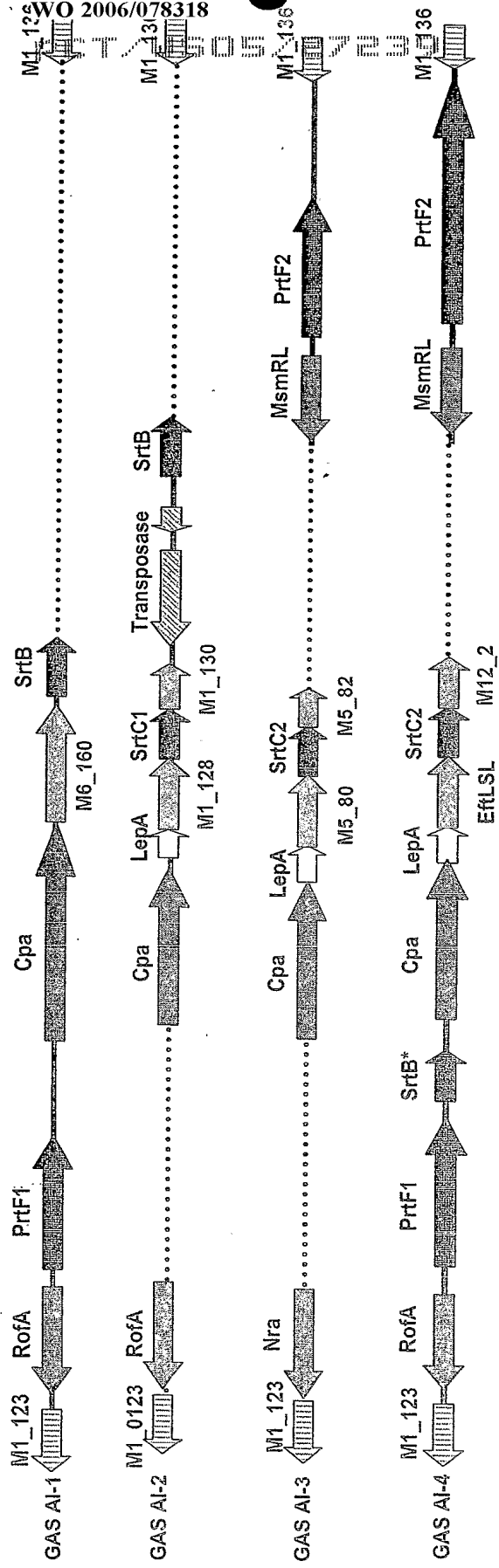
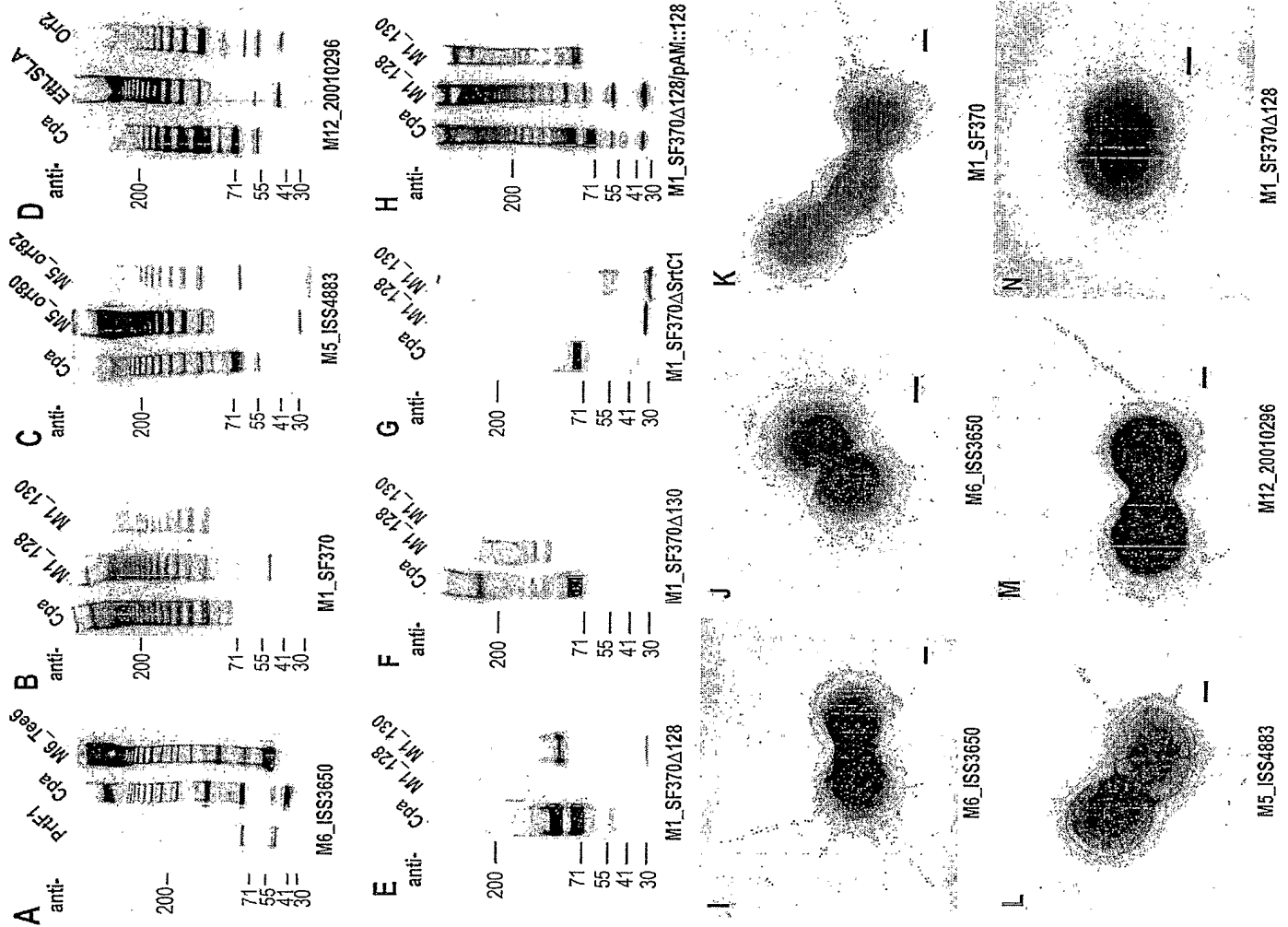


FIGURE 162

Figure 163



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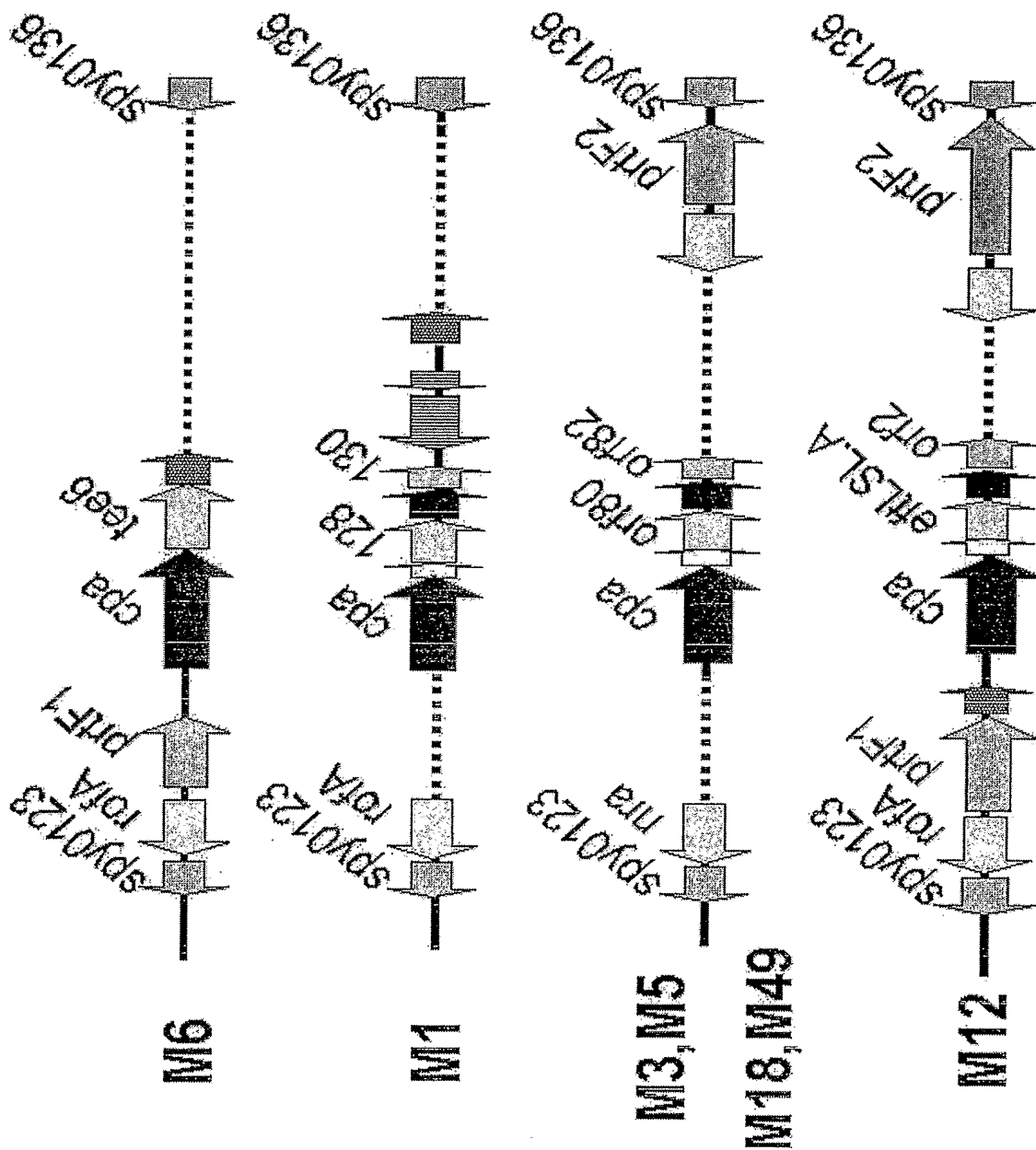


Figure 164

PCT/US05/27239363/487

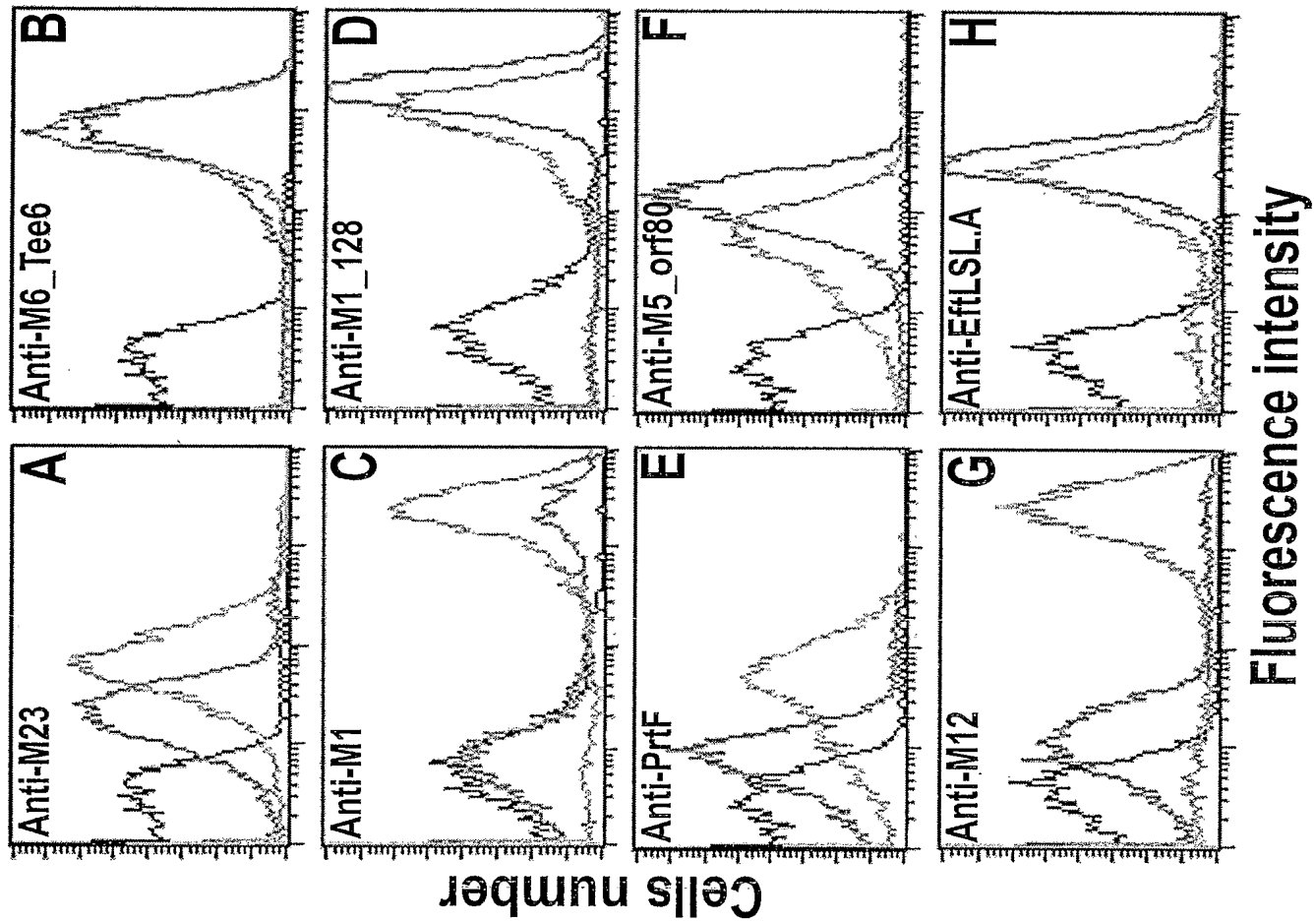


Figure 165

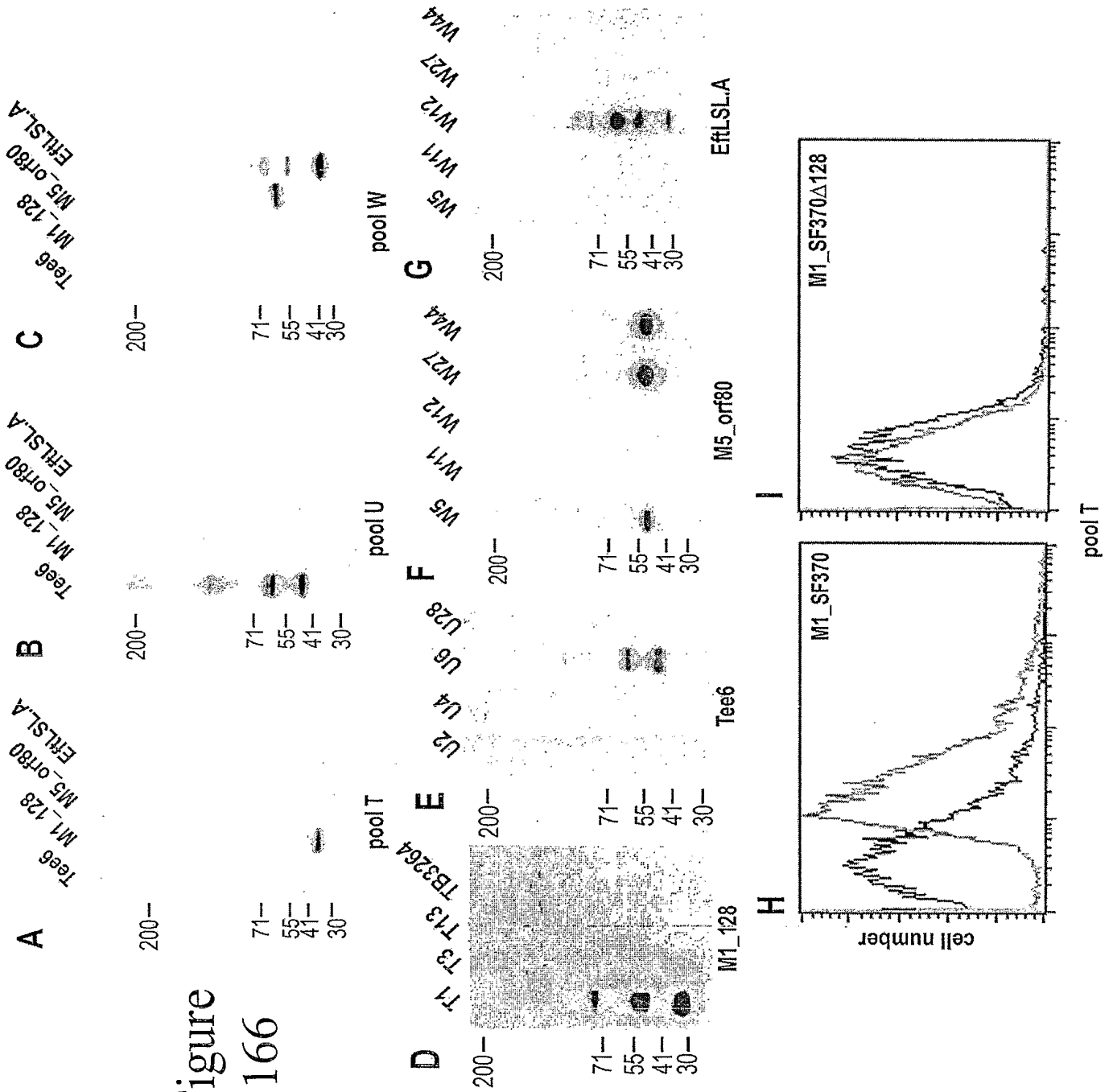


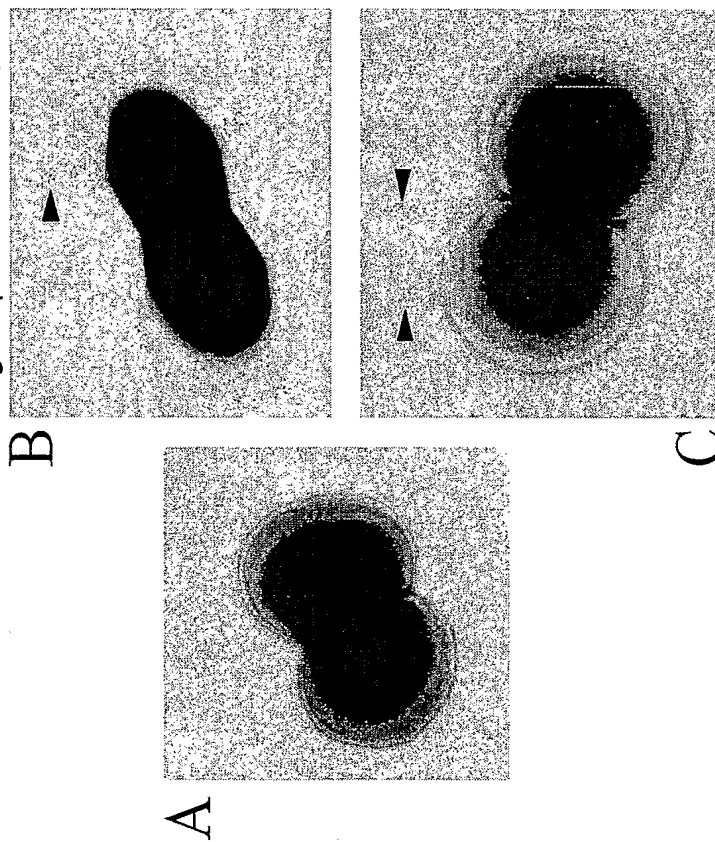
Figure 167

Strain	M-type	PCR					AI	Sequence
		SrtB	SrtC1	SrtC2	MsmRL	SipA2		
2724	6	+	-	-	-	-	1	
2894	6	+	-	-	-	-	1	
3650	6	+	-	-	-	-	1	
5529	6	+	-	-	-	-	1	
Dsm2071	23	+	-	-	-	-	1	+
SF370	1	+	+	-	-	-	2	literature
2580	1	+	+	-	-	-	2	
2913	1	+	+	-	-	-	2	
3280	1	+	+	-	-	-	2	
3348	1	+	+	-	-	-	2	
2719	?	+	+	-	-	-	2	
2721	3	-	-	+	+	+	3	
3040	3	-	-	+	+	+	3	
3135	3	-	-	+	+	+	3	
3776	44 ?	-	-	+	+	+	3	+
4959	77	-	-	+	+	+	3	+
4088	Clinical isolate	-	-	+	+	+	3	
2728	12	+	-	+	+	+	4	
2720	9	+	-	+	+	+	4	+
2727	11	+	-	+	+	+	4	+
4436	28	+	-	+	+	+	4	+
5481	44 ?	+	-	+	+	+	4	+
4538	50	+	-	+	+	+	4	+
3789	78	+	-	+	+	+	4	+
4883	5	+	-	+	+	+	4	
5476	89	+	-	+	+	+	4	
5495	?	+	-	+	+	+	4	
2722	4	-	-	-	-	-	?	
2723	5?	-	-	-	-	-	?	
2725	8	-	-	+	-	-	?	
2726	2	-	-	-	-	-	?	
2634	4	-	-	-	-	-	?	
5531	75	+	+	-	-	-	?	In progress

Figure 168

Immuno-electronmicroscopy

(Immunogold Negative Staining,
1° α - 80, 2° α - mouse gold particles 10nm)



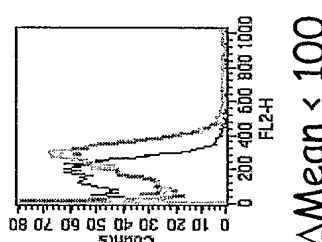
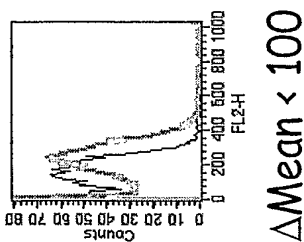
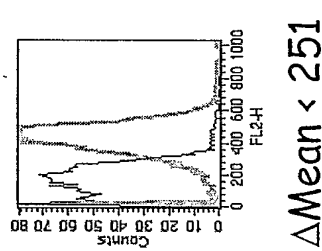
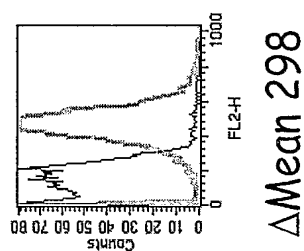
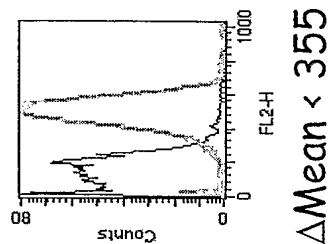
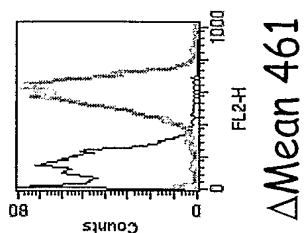
<i>L.lactis</i>	<i>L.lactis</i> + AI-1
-	+

Figure 169

GBS JM9130013

L. lactis + AI-1

L. lactis

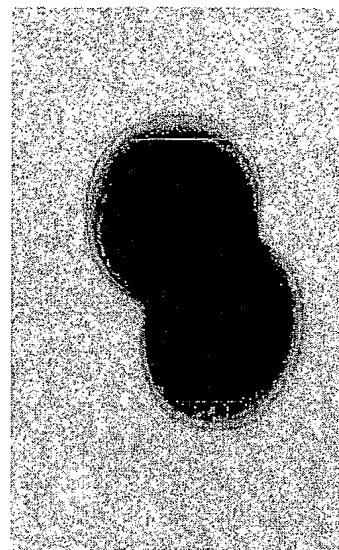


α -80

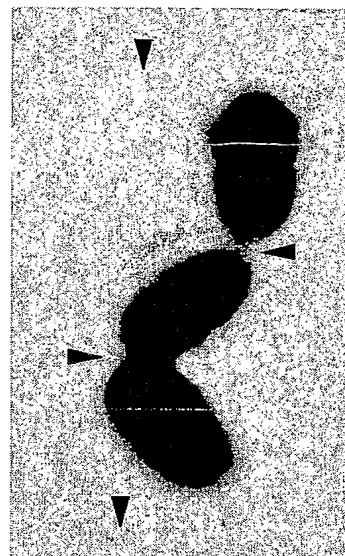
α -104

Figure 170

Phase contrast Microscopy **Immuno-electronmicroscopy**
 (Immunogold Negative Staining,
 1° α -80, 2° α -mouse gold particles 10nm)



L. lactis

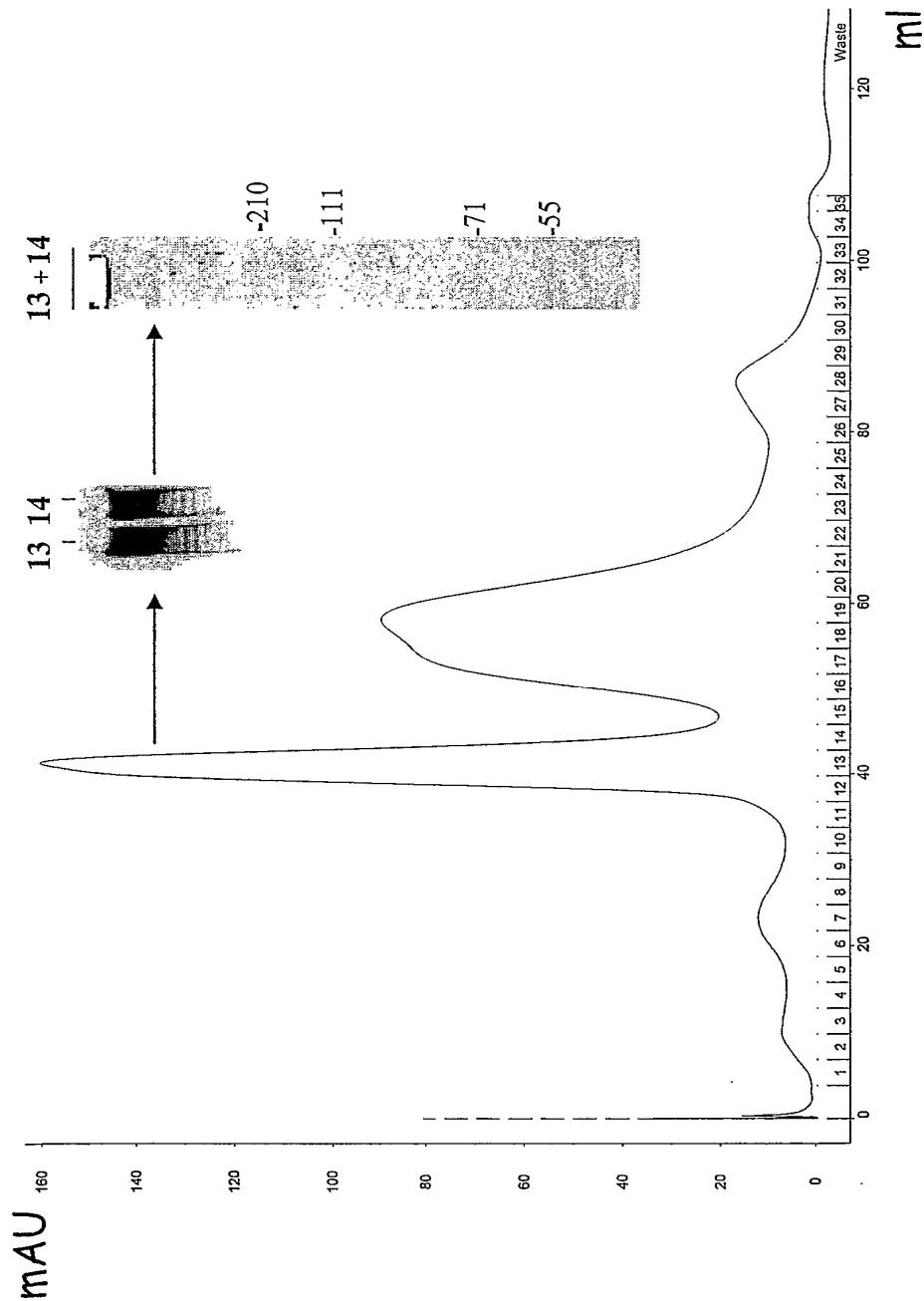


L. lactis + AI-1

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Figure 171

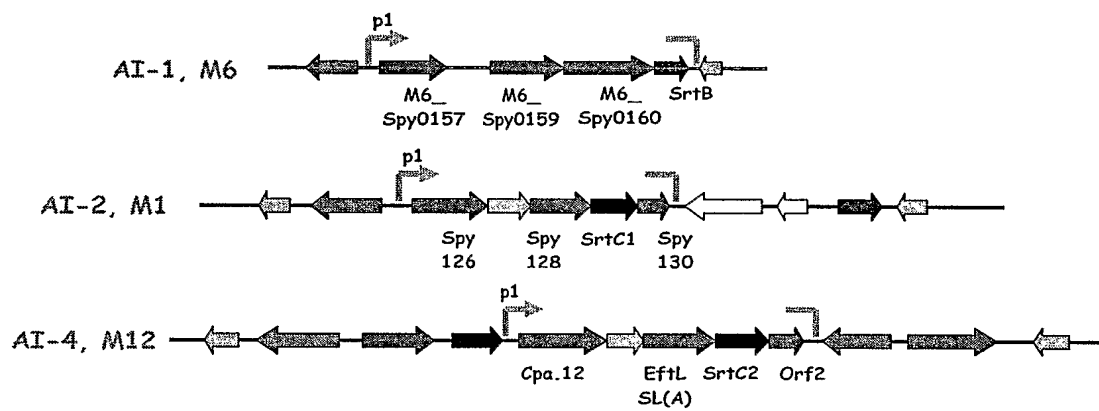
Gel filtration on Sepharyl HR 400

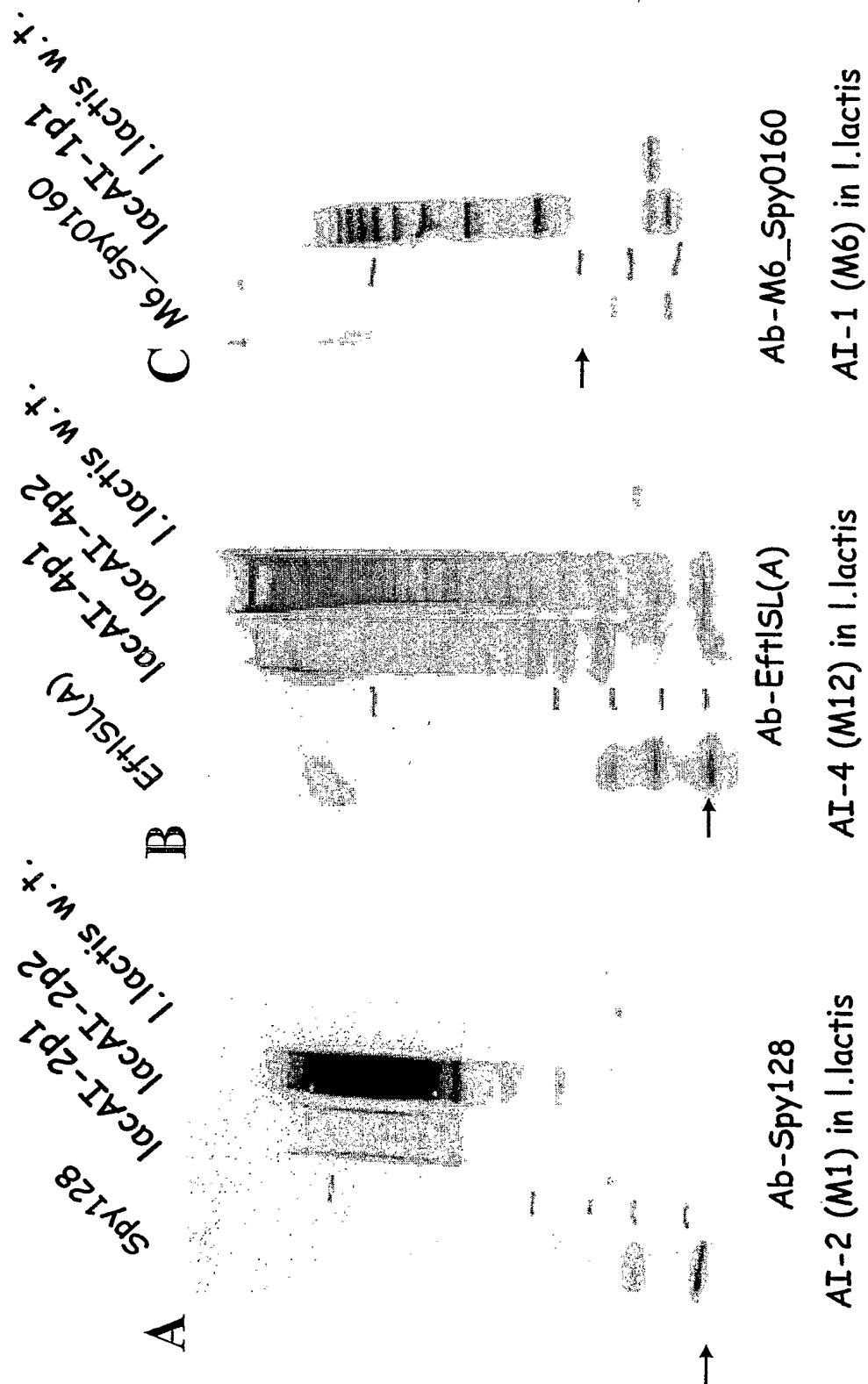


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Figure 172

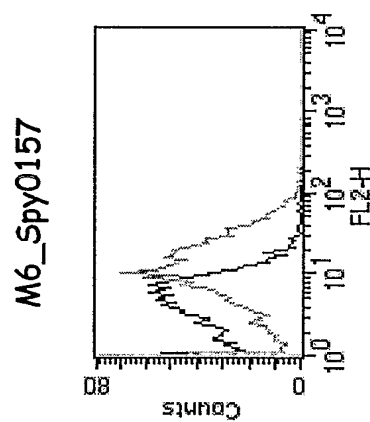




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Figure 174



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Figure 175

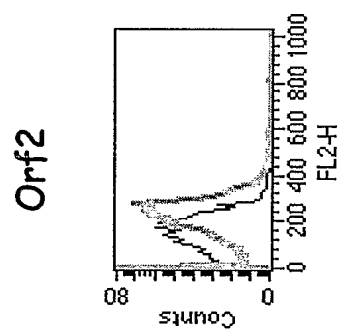
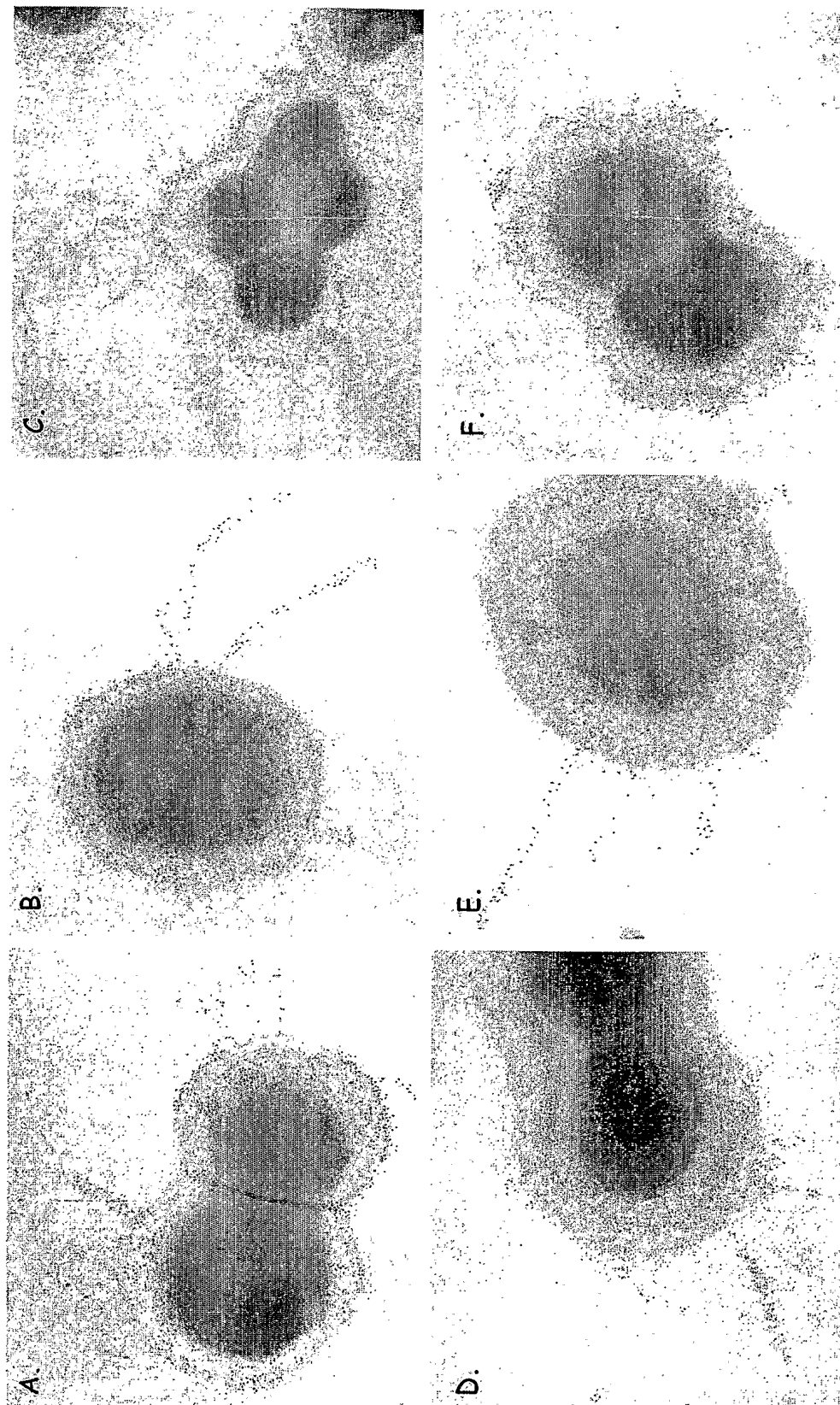


Figure 176



Immunogold labeling with antibodies against: A. B. C. D. E. M6_Spy0160; F. M6_Spy0159

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Figure 177

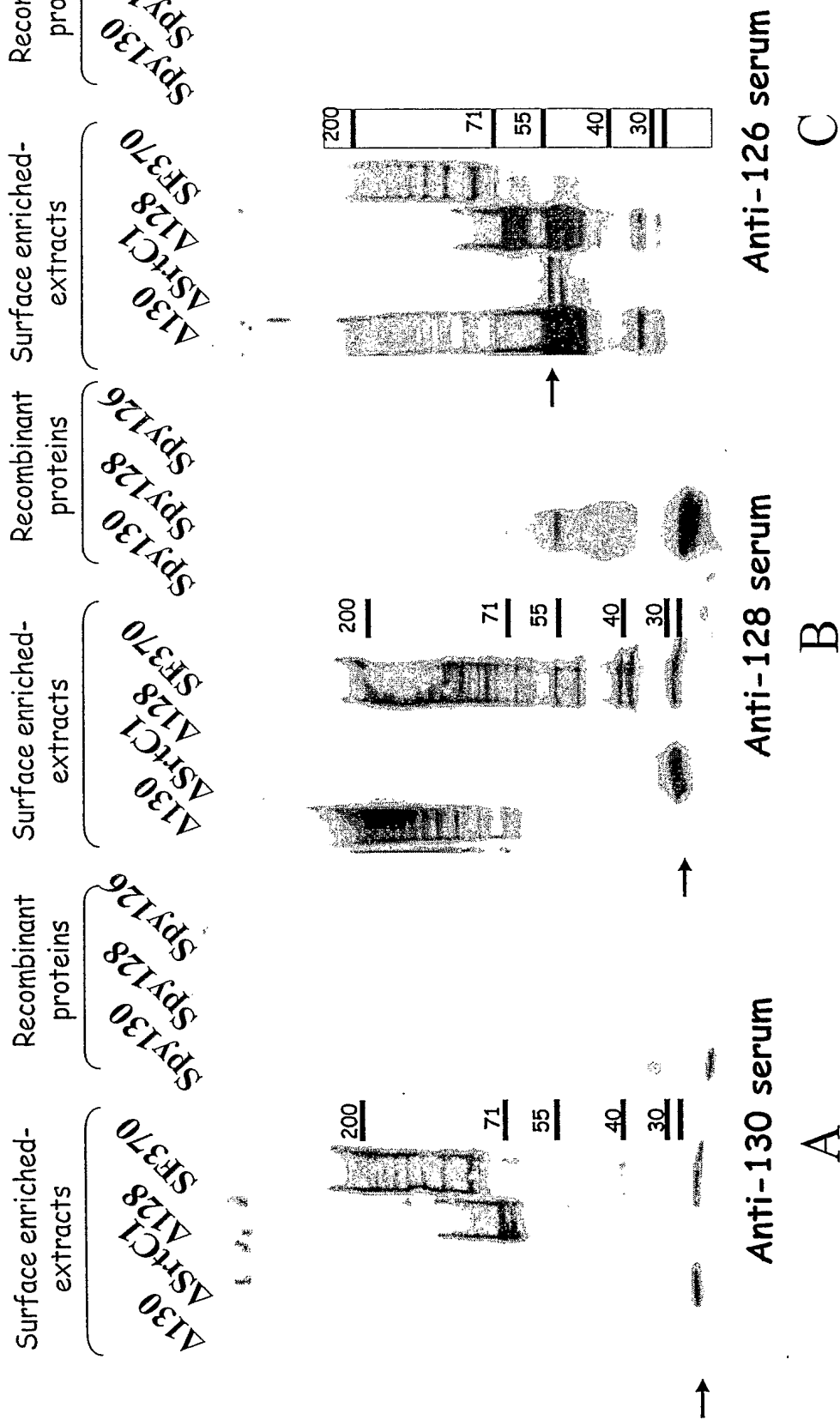
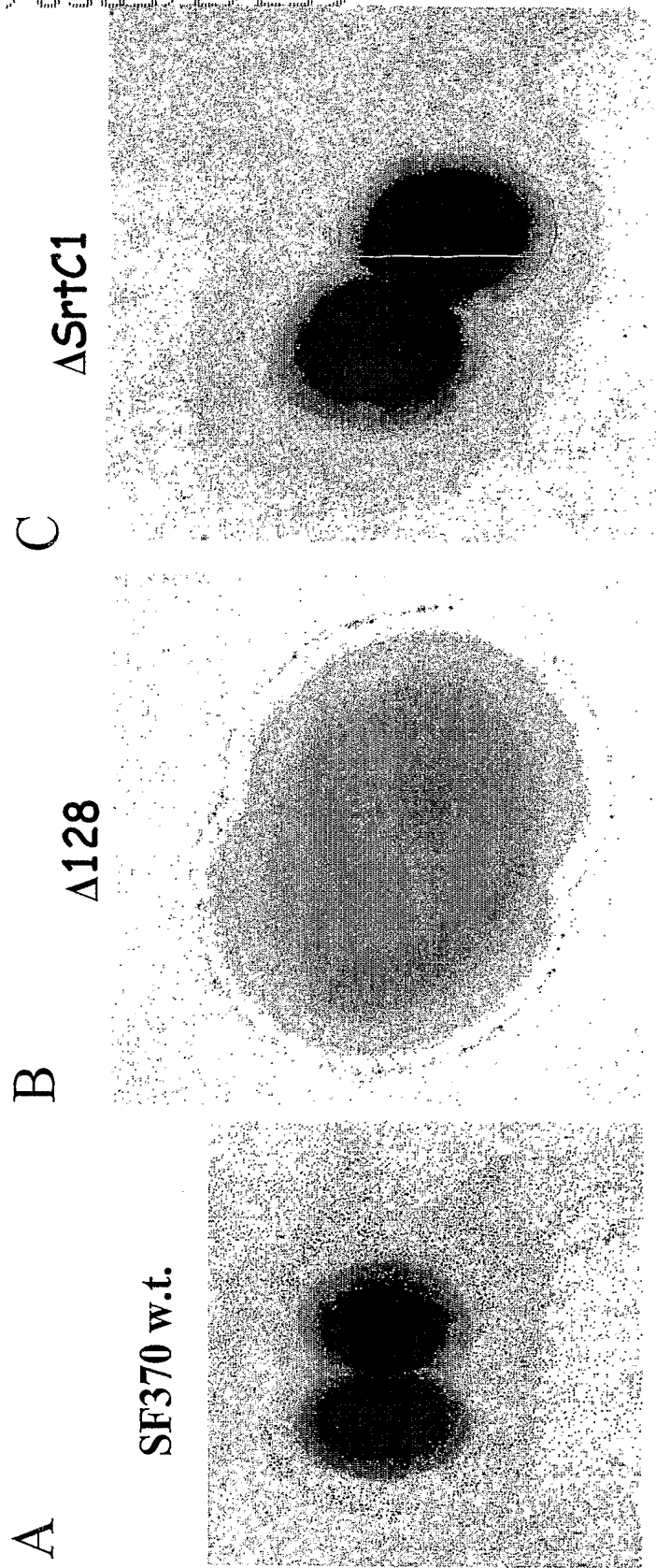


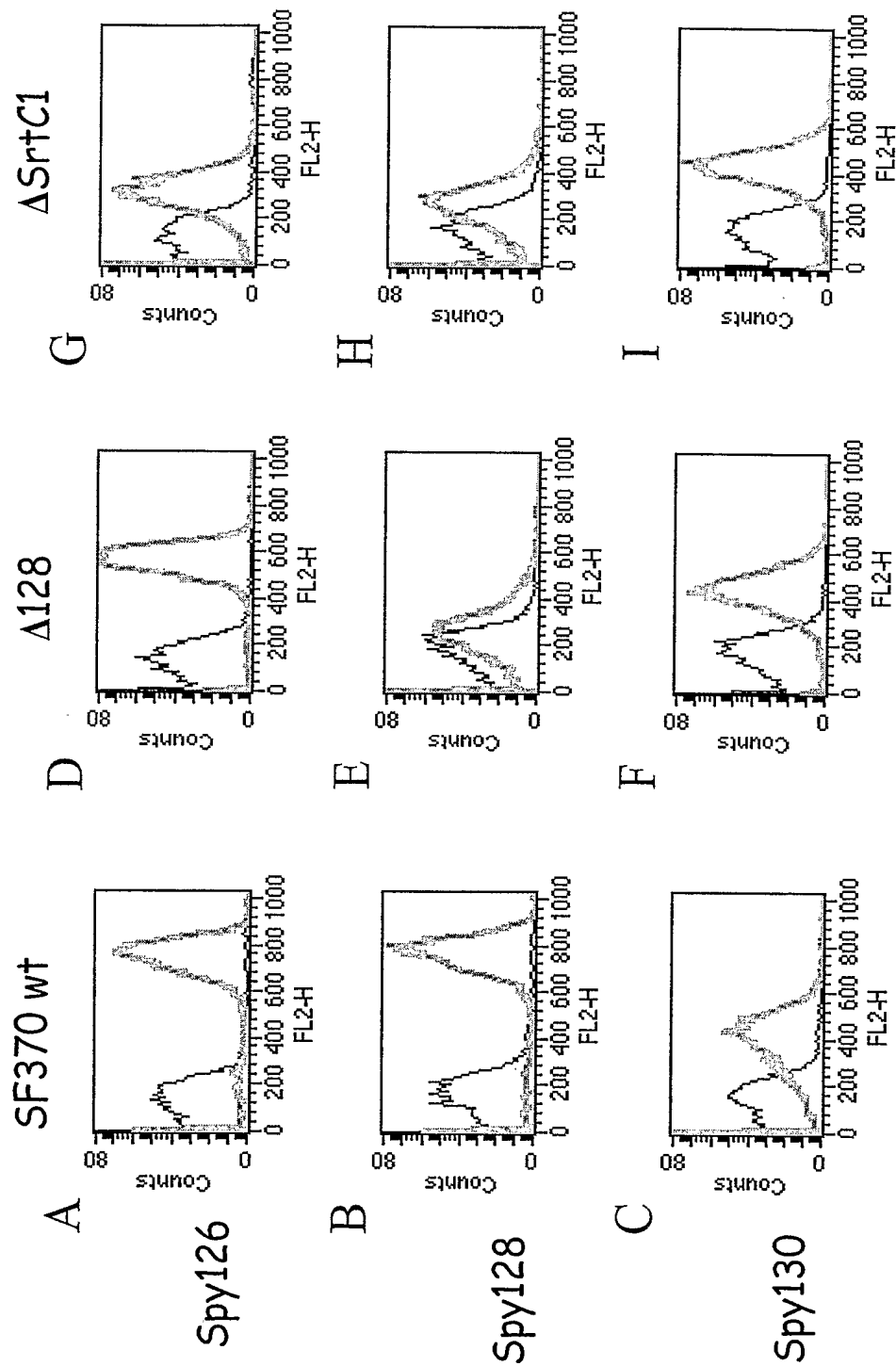
Figure 178



Immuno-gold labeling with sera against Spy128

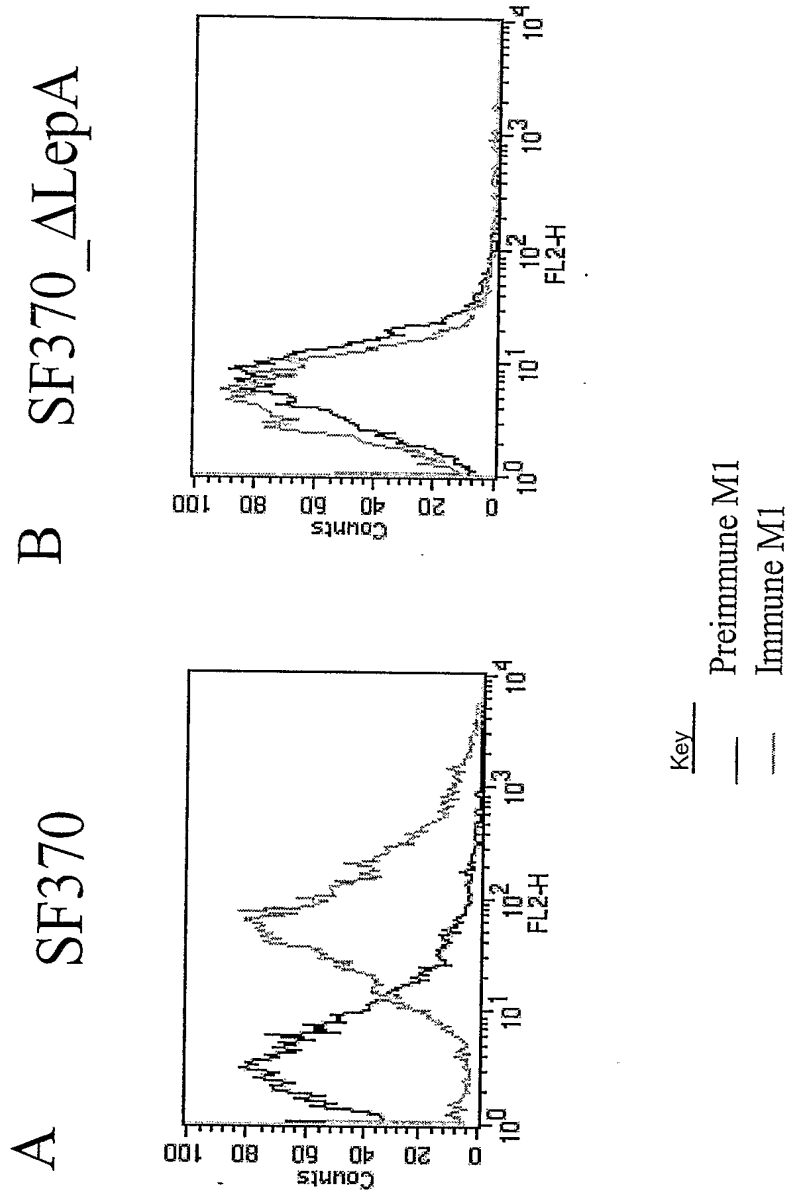
Comparison of wild type and mutant strain by Immunoelectron Microscopy show that Spy128- or SrtC1-lacking bacteria are not able to assemble pili. SrtC1, therefore, is absolutely required for pilus assembly but not for surface anchoring.

Figure 179



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Figure 180



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Figure 181

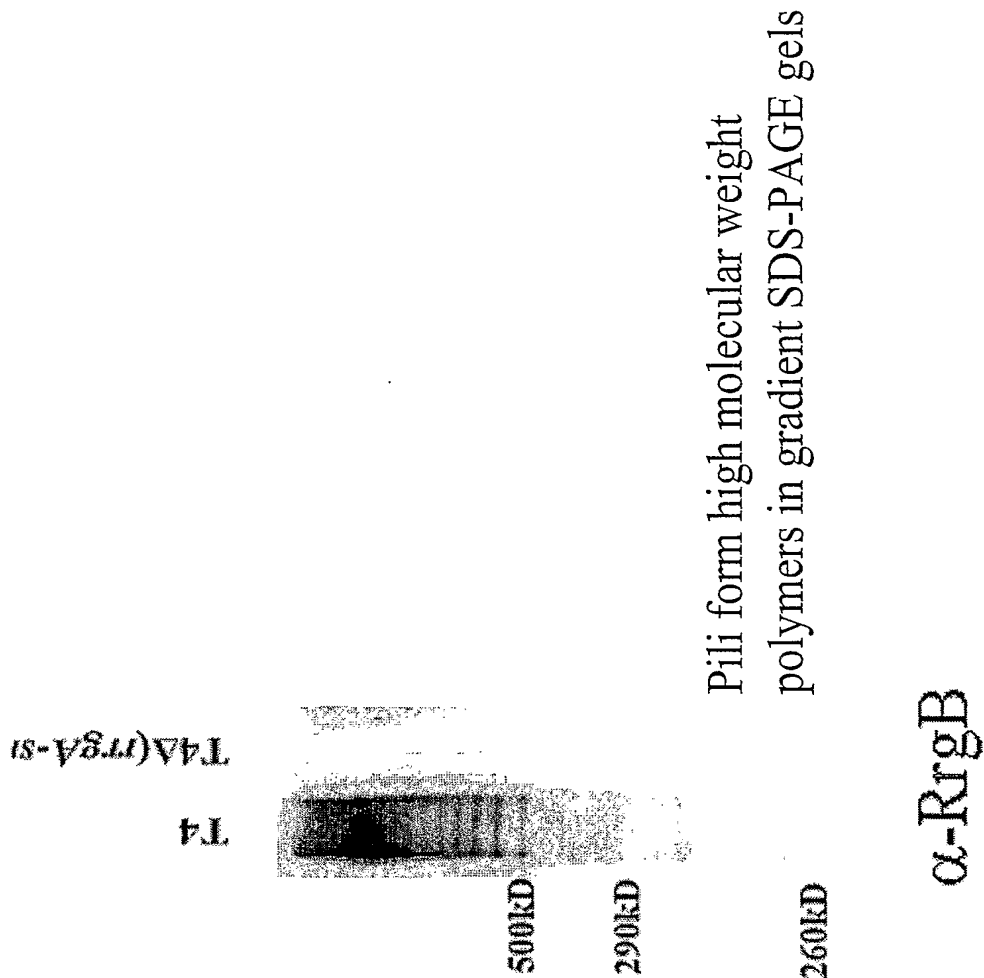
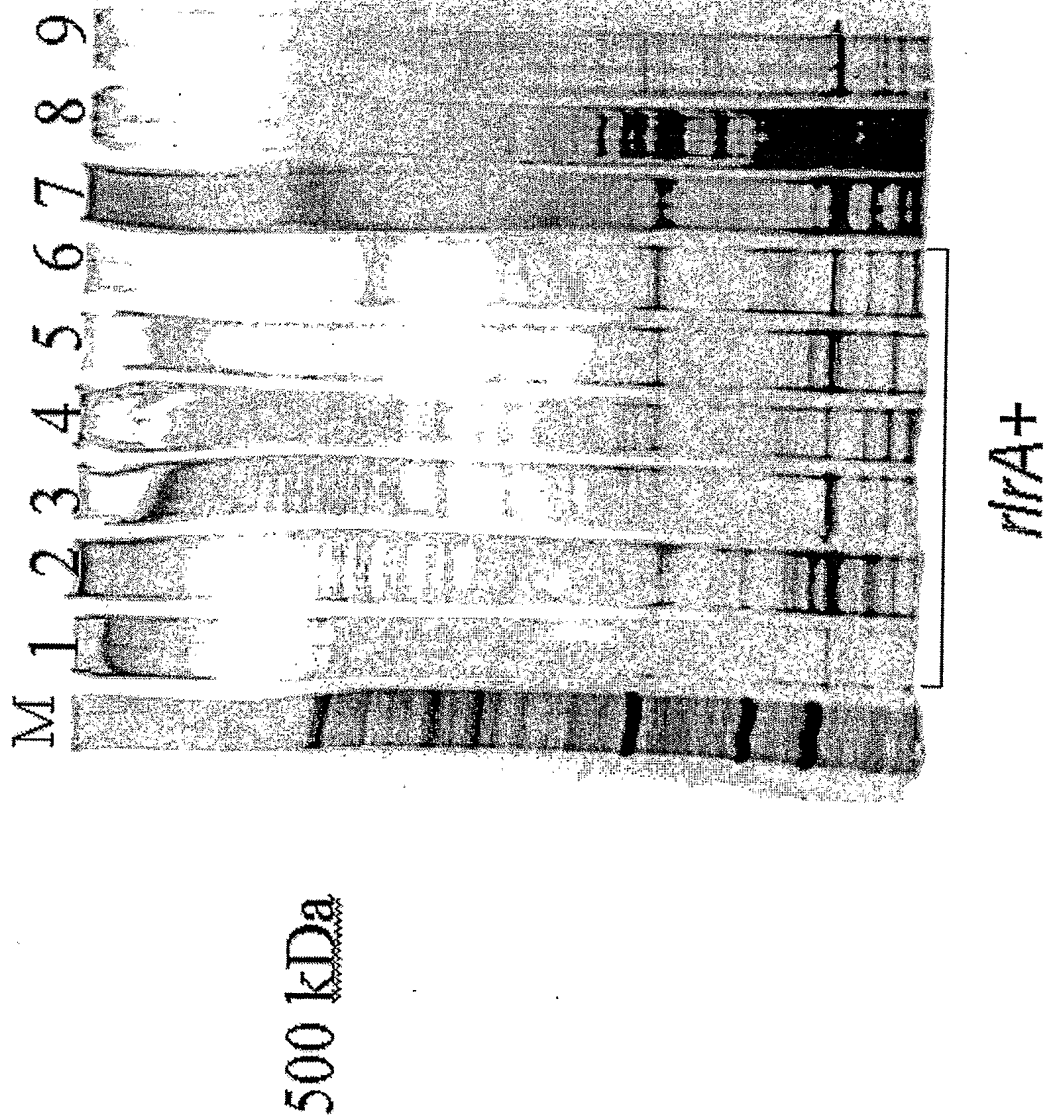


Figure 182



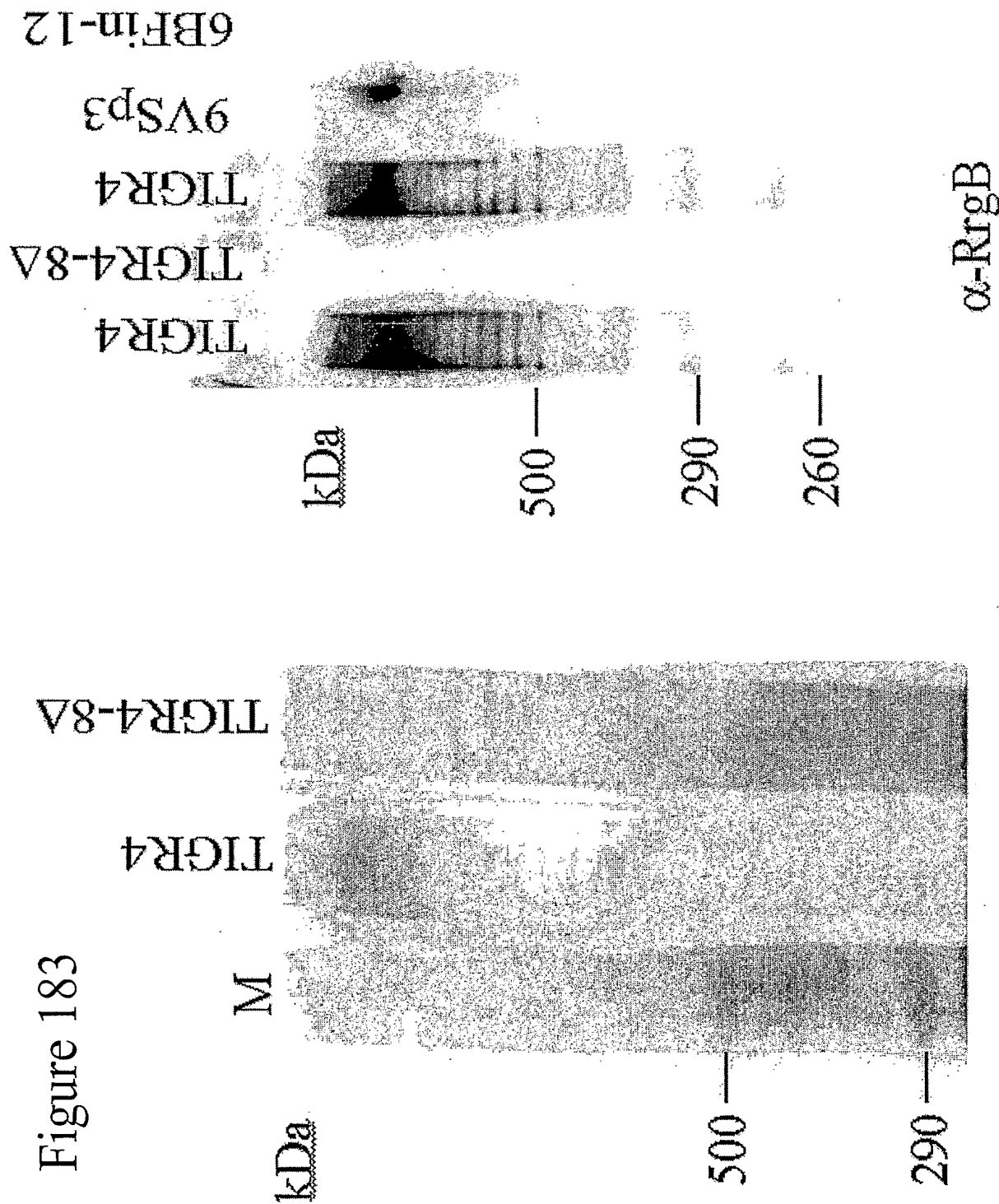


Figure 183

Silver stained gel 3-8%

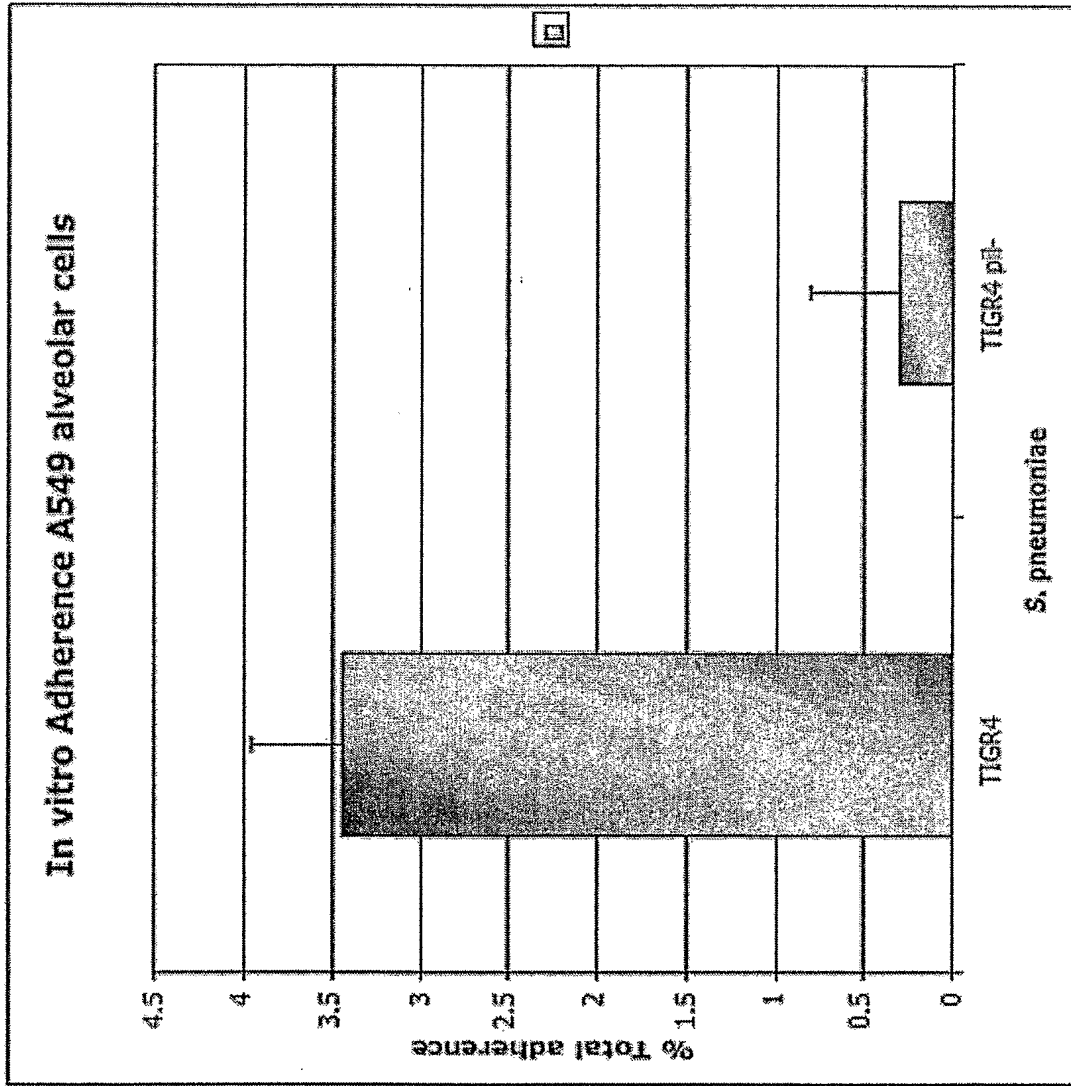
α -RrgB

Anti-RrgB TIGR4 recognized the 9v pili

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Figure 184



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Figure 185



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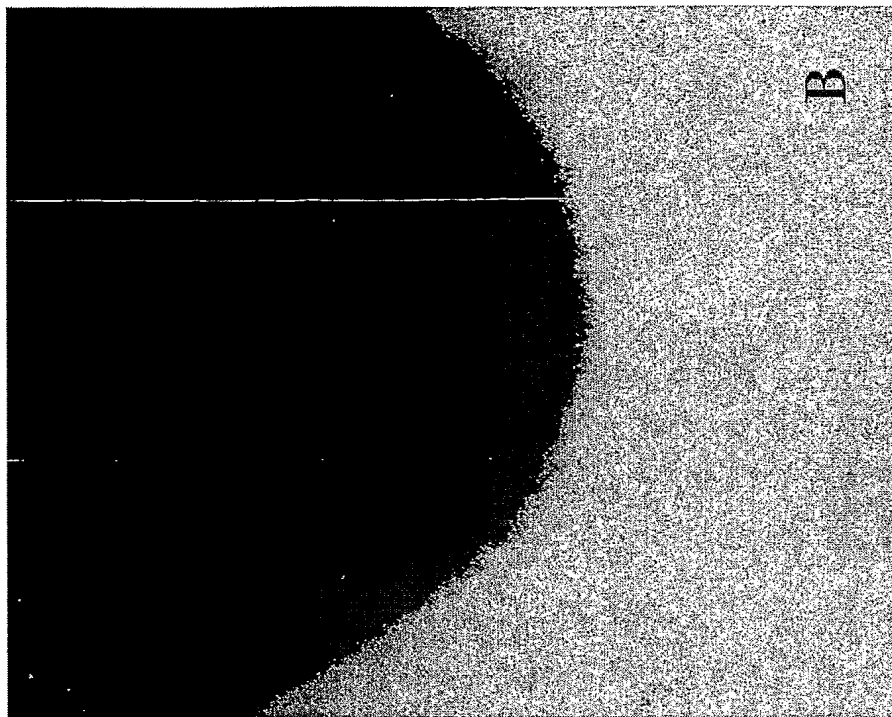


Figure 186

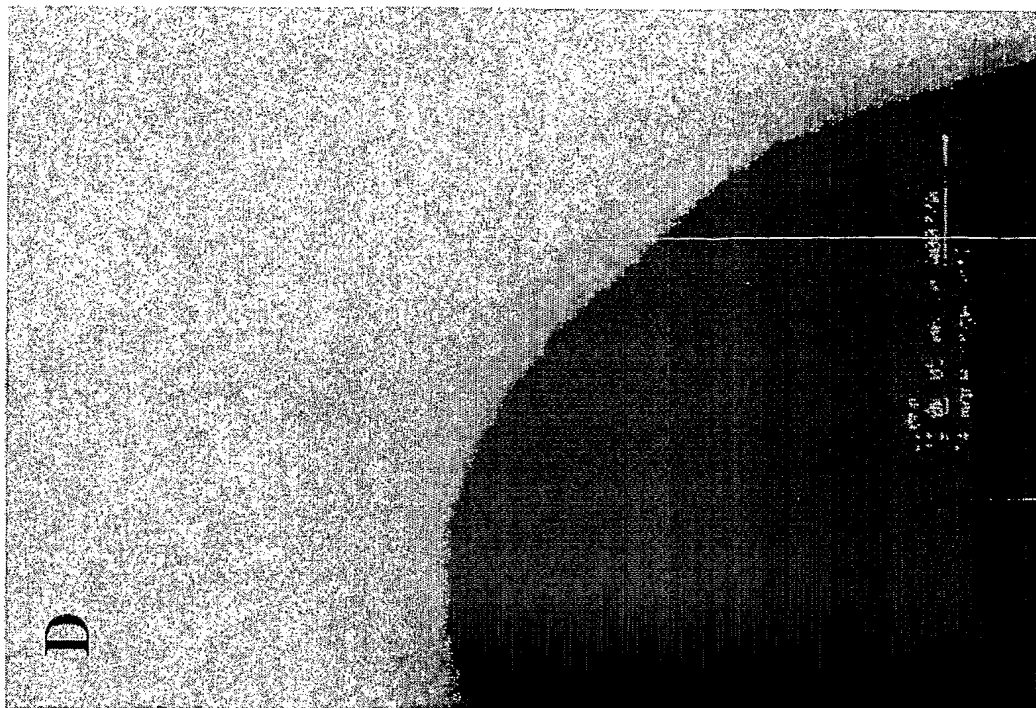


Figure 188

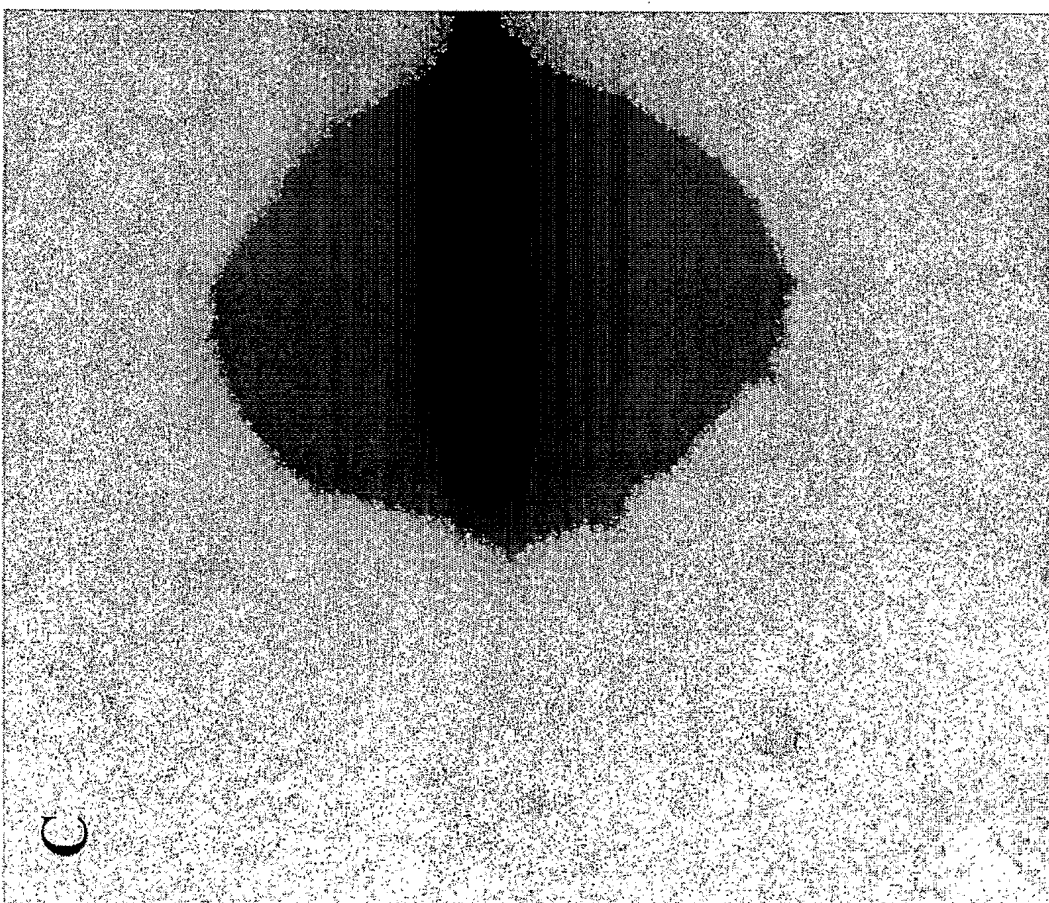


Figure 187

Figure 189

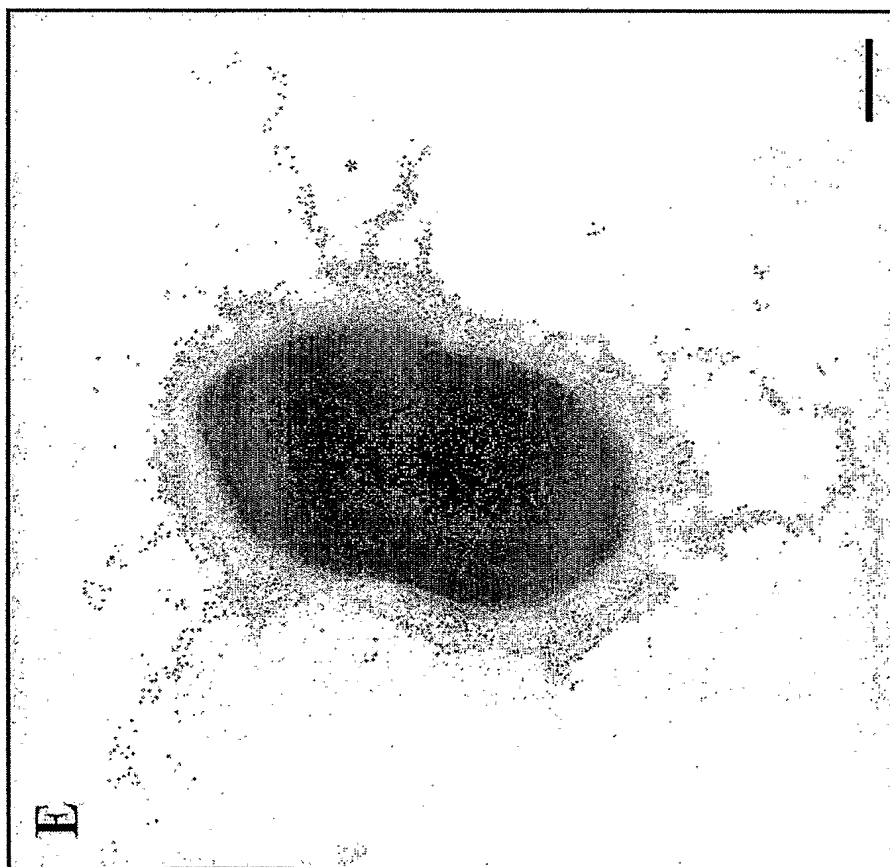


Figure 190

S. pneumoniae pili proteins: sp0462 (Rrg.A)

Expression and purification:

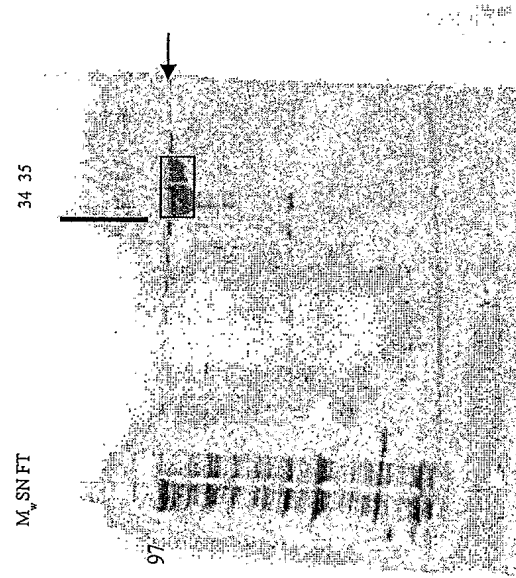
- pET 21b+-*rrg.A-6*
- purified in soluble form (stored at -80°C ; in $\text{NaCl}_{\text{physiol.}}$)



Results:

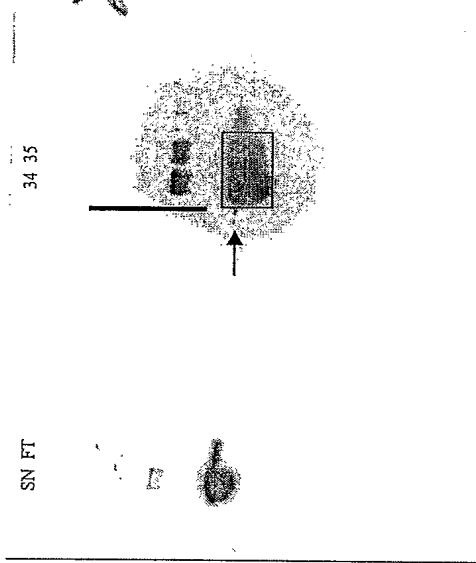
- protein conc.: 1,1 mg/ml

A



SDS-page

B



Western blot (anti-HIS)

S. pneumoniae pili proteins – antibody production (mice)

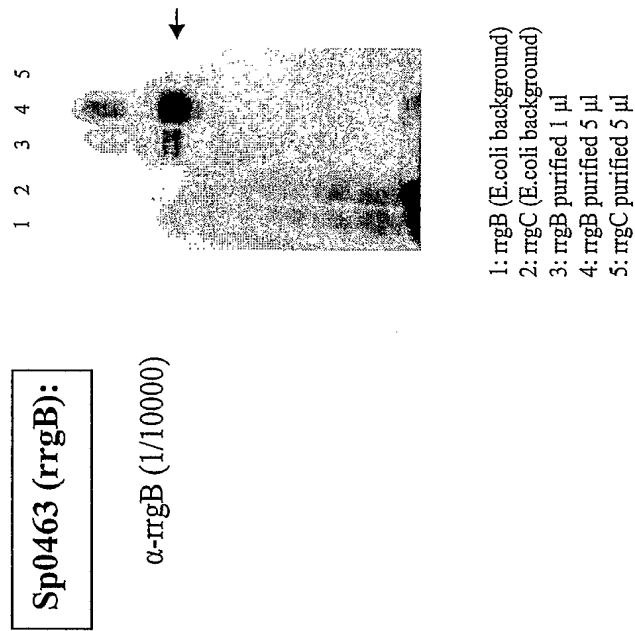


Figure 191

S. pneumoniae pili proteins – antibody production (mice)

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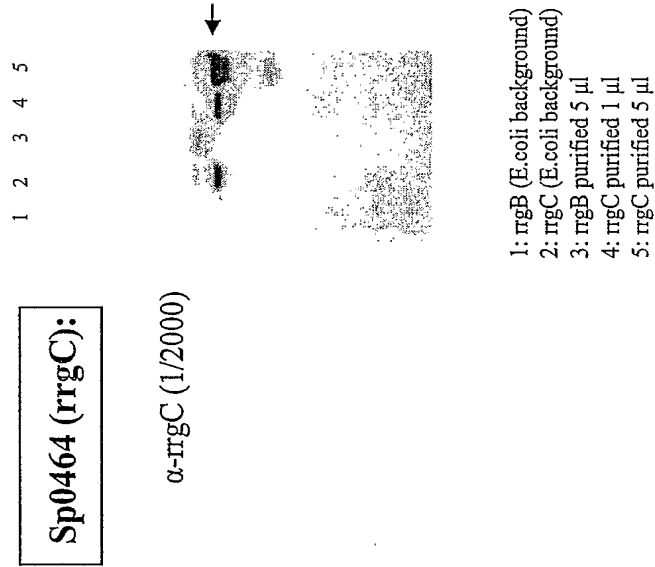
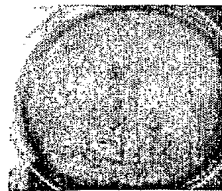


Figure 192

S. pneumoniae TIGR4 pilus purification I – cultivation + digestion



S. pneumoniae TIGR4
Blood plates
ON/37°C/13h

- Resuspension in PBS/washing
- Resuspension in PPB (4-6 plates/ml)
(20% sucrose, 10mM MgCl₂,
50mM NaPPi pH6.3)

- Digestion with Mutanolysin
(N-Acetyl Muramidase)
37°C, ~10 h

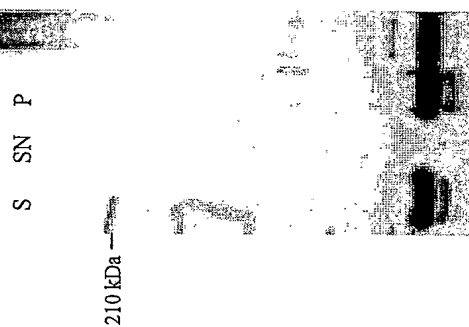
Pellet
SN

Sucrose Density
gradient centrifugation

T4

D39

T4/S



Western (1.AK. α-irgB)

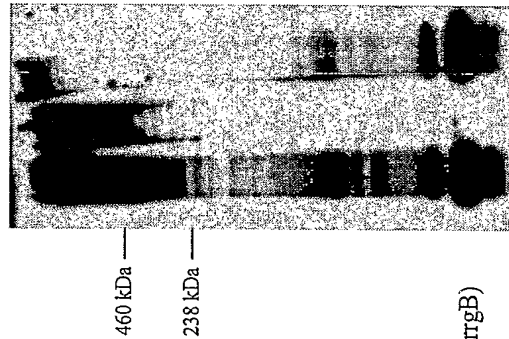


Figure 193

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S. pneumoniae TIGR4 pilus purification II - Sucrose density gradient centrifugation

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950µl SN
25-56% linear sucrose gradient
SW40; 38000, 4°C, 16h



24 x 500 µl fractions
(Gradient master)



Gel filtration

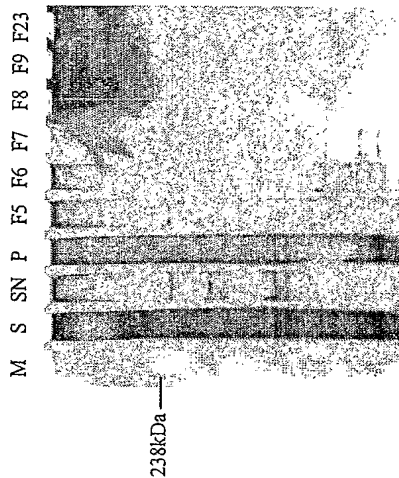
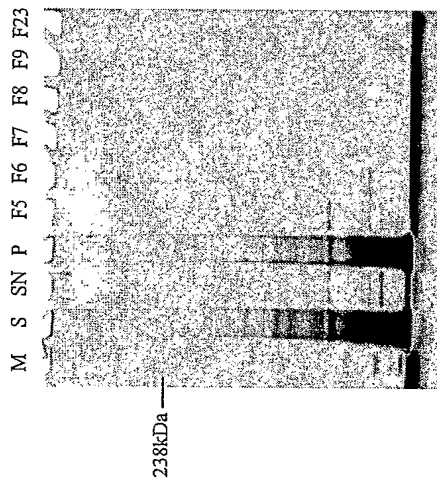
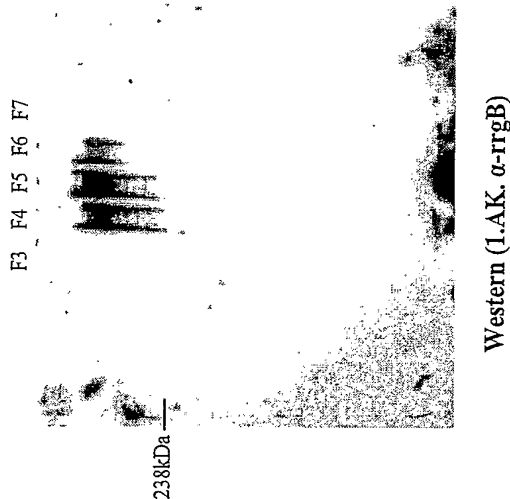
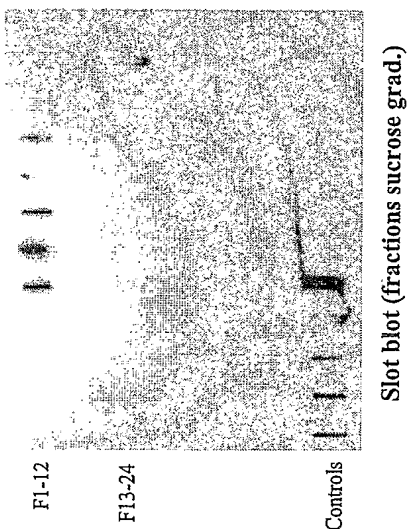


Figure 194

400 μ l Fr.5
Superdex 200

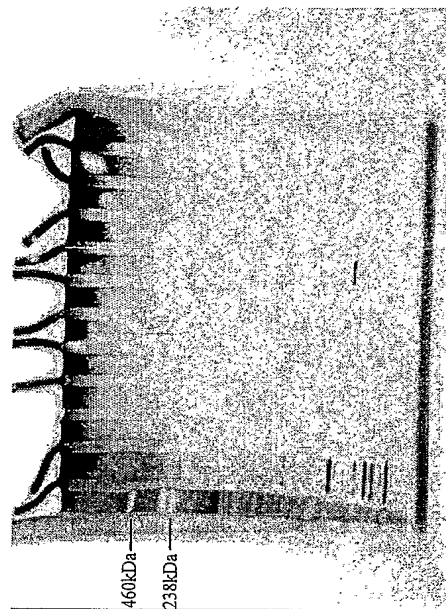
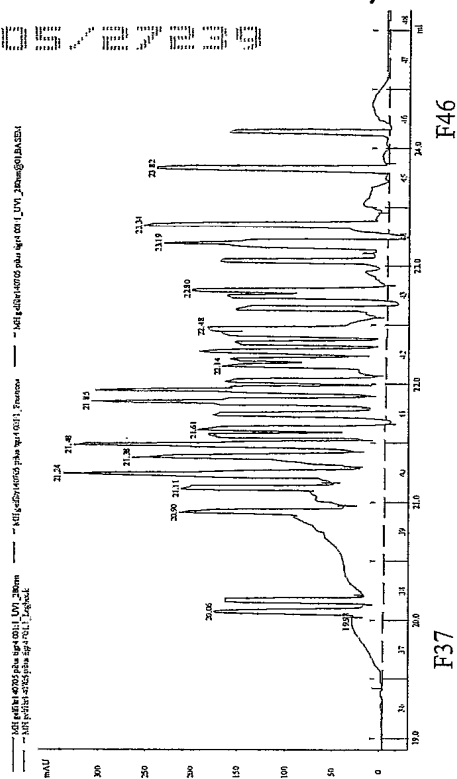
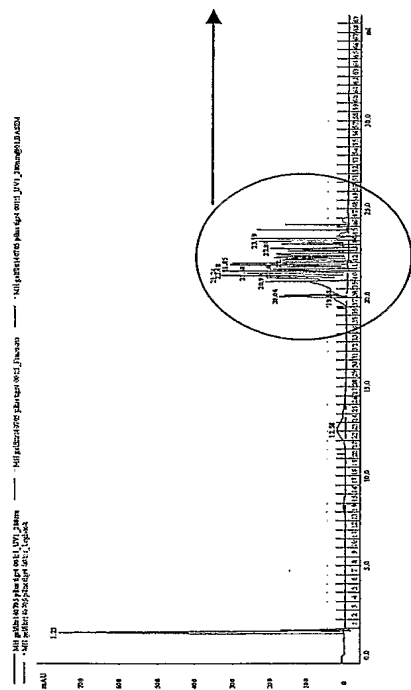


Figure 195

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14CSR -----GTTTAGGCGCTTTTCATTATAAGTCTTATGGGACTTTTTTGATACTCAAAAAGC
670 TGAGTTGTTTAGGCGCTTTTCATTATAAGTCTTATGGGACTTTTTTGATACTCAAAAAGC
6BF -----GTTTAGGCGCTTTTCATTATAAGTCTTATGGGACTTTTTTGATACTCAAAAAGC
6BSP -----GCGCTTTTCATTATAAGTCTTATGGGACTTTTTTGATACTCAAAAAGC
19AH -----GTTTAGGCGCTTTTCATTATAAGTCTTATGGGACTTTTTTGATACTCAAAAAGC
23FPO -----TTAGGCGCTTTTCATTATAAGTCTTATGGGACTTTTTTGATACTCAAAAAGC
19FTW -----TTTTTCATTATAAATCTTATGGGACTTTTTTGATACTCAAAAAGC
9VSP -----TTAGGCGCTTTTCATTATAAGTCTTATGGGACTTTTTTGATACTCAAAAAGC
TIGR4 -----TTAGGCGCTTTTCATTATAAGTCTTATGGGACTTTTTTGATACTCAAAAAGC
23FTW -----GCGCTTTTCATTATAAGTCTTATGGGACTTTTTTGATACTCAAAAAGC

14CSR CCTATAATCTCCACAGTGGGATTTACCCACTACAGAAATTATAGAGCCAGAAAAACACT
670 CCTATAATCTCCACAGTGGGATTTACCCACTACAGAAATTATAGAGCCAGAAAAACACT
6BF CCTATAATCTCCACAGTGGGATTTACCCACTACAGAAATTATAGAGCCAGAAAAACACT
6BSP CCTATAATCTCCACAGTGGGATTTACCCACTACAGAAATTATAGAGCCAGAAAAACACT
19AH CCTATAATCTCCACAGTGGGATTTACCCACTACAGAAATTATAGAGCCAGAAAAACACT
23FPO CCTATAATCTCCACAGTGGGATTTACCCACTACAGAAATTATAGAGCCAGAAAAACACT
19FTW CCTATAATCTCCACAGTGGGATTTACCCACTACAGAAATTATAGAGCCAGAAAAACACT
9VSP CCTATAATCTCCACAGTGGGATTTACCCACTACAGAAATTATAGAGCCAGAAAAACACT
TIGR4 CCTATAATCTCCACAGTGGGATTTACCCACTACAGAAATTATAGAGCCAGAAAAACACT
23FTW CCTATAATCTCCACAGTGGGATTTACCCACTACAGAAATTATAGAGCCAGAAAAACACT

14CSR TTTGTTCACTAGCAGAACTAGAGAGCAGAAGTGTTTTCTGTTTCTGTTTACCCAAAAC
670 TTTGTTCACTAGCAGAACTAGAGAGCAGAAGTGTTTTCTGTTTCTGTTTACCCAAAAC
6BF TTTGTTCACTAGCAGAACTAGAGAGCAGAAGTGTTTTCTGTTTCTGTTTACCCAAAAC
6BSP TTTGTTCACTAGCAGAACTAGAGAGCAGAAGTGTTTTCTGTTTCTGTTTACCCAAAAC
19AH TTTGTTCACTAGCAGAACTAGAGAGCAGAAGTGTTTTCTGTTTCTGTTTACCCAAAAC
23FPO TTTGTTCACTAGCAGAACTAGAGAGCAGAAGTGTTTTCTGTTTCTGTTTACCCAAAAC
19FTW TTTGTTCACTAGCAGAACTAGAGAGCAGAAGTGTTTTCTGTTTCTGTTTACCCAAAAC
9VSP TTTGTTCACTAGCAGAACTAGAGAGCAGAAGTGTTTTCTGTTTCTGTTTACCCAAAAC
TIGR4 TTTGTTCACTAGCAGAACTAGAGAGCAGAAGTGTTTTCTGTTTCTGTTTACCCAAAAC
23FTW TTTGTTCACTAGCAGAACTAGAGAGCAGAAGTGTTTTCTGTTTCTGTTTACCCAAAAC

14CSR TGGGAAATATGGGGATAAGAATAGAGATGGCTTAGGAAGCCCCCTTTTGTGTGTAGACAG
670 TGGGAAATATGGGGATAAGAATAGAGATGGCTTAGGAAGCCCCCTTTTGTGTGTAGACAG
6BF TGGGAAATATGGGGATAAGAATAGAGATGGCTTAGGAAGCCCCCTTTTGTGTGTAGACAG
6BSP TGGGAAATATGGGGATAAGAATAGAGATGGCTTAGGAAGCCCCCTTTTGTGTGTAGACAG
19AH TGGGAAATATGGGGATAAGAATAGAGATGGCTTAGGAAGCCCCCTTTTGTGTGTAGACAG
23FPO TGGGAAATATGGGGATAAGAATAGAGATGGCTTAGGAAGCCCCCTTTTGTGTGTAGACAG
19FTW TGGGAAATATGGGGATAAGAATAGAGATGGCTTAGGAAGCCCCCTTTTGTGTGTAGACAG
9VSP TGGGAAATATGGGGATAAGAATAGAGATGGCTTAGGAAGCCCCCTTTTGTGTGTAGACAG
TIGR4 TGGGAAATATGGGGATAAGAATAGAGATGGCTTAGGAAGCCCCCTTTTGTGTGTAGACAG
23FTW TGGGAAATATGGGGATAAGAATAGAGATGGCTTAGGAAGCCCCCTTTTGTGTGTAGACAG

14CSR TACGATGAACCTATAACAAATAGTGAGCCTTTTGTAGCAATCATTCGACCCGTTTGTCAA
670 TACGATGAACCTATAACAAATAGTGAGCCTTTTGTAGCAATCATTCGACCCGTTTGTCAA
6BF TACGATGAACCTATAACAAATAGTGAGCCTTTTGTAGCAATCATTCGACCCGTTTGTCAA
6BSP TACGATGAACCTATAACAAATAGTGAGCCTTTTGTAGCAATCATTCGACCCGTTTGTCAA
19AH TACGATGAACCTATAACAAATAGTGAGCCTTTTGTAGCAATCATTCGACCCGTTTGTCAA
23FPO TACGATGAACCTATAACAAATAGTGAGCCTTTTGTAGCAATCATTCGACCCGTTTGTCAA
19FTW TACGATGAACCTATAACAAATAGTGAGCCTTTTGTAGCAATCATTCGACCCGTTTGTCAA
9VSP TACGATGAACCTATAACAAATAGTGAGCCTTTTGTAGCAATCATTCGACCCGTTTGTCAA
TIGR4 TACGATGAACCTATAACAAATAGTGAGCCTTTTGTAGCAATCATTCGACCCGTTTGTCAA
23FTW TACGATGAACCTATAACAAATAGTGAGCCTTTTGTAGCAATCATTCGACCCGTTTGTCAA

Figure 196A

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14CSR AAGCCTCTTTTCGGATATCTACAATTGTCTGATAGATGAGACGCTGTTGGCTAACATGCA
670 AAGCCTCTTTTCGGATATCTACAATTGTCTGATAGATGAGACGCTGTTGGCTAACATGCA
6BF AAGCCTCTTTTCGGATATCTACAATTGTCTGATAGATGAGACGCTGTTGGCTAACATGCA
6BSP AAGCCTCTTTTCGGATATCTACAATTGTCTGATAGATGAGACGCTGTTGGCTAACATGCA
19AH AAGCCTCTTTTCGGATATCTACAATTGTCTGATAGATGAGACGCTGTTGGCTAACATGCA
23FPO AAGCCTCTTTTCGGATATCTACAATTGTCTGATAGATGAGACGCTGTTGGCTAACATGCA
19FTW AAGCCTCTTTTCGGATATCTACAATTGTCTGATAGATGAGACGCTGTTGGCTAACATGCA
9VSP AAGCCTCTTTTCGGATATCTACAATTGTCTGATAGATGAGACGCTGTTGGCTAACATGCA
TIGR4 AAGCCTCTTTTCGGATATCTACAATTGTCTGATAGATGAGACGCTGTTGGCTAACATGCA
23FTW AAGCCTCTTTTCGGATATCTACAATTGTCTGATAGATGAGACGCTGTTGGCTAACATGCA

14CSR AATCTAAGGCAATCGTCAAAAAGTGATGTTTCCCTTTGGGATACTGCTTTTTAACGTAAG
670 AATCTAAGGCAATCGTCAAAAAGTGATGTTTCCCTTTGGGATACTGCTTTTTAACGTAAG
6BF AATCTAAGGCAATCGTCAAAAAGTGATGTTTCCCTTTGGGATACTGCTTTTTAACGTAAG
6BSP AATCTAAGGCAATCGTCAAAAAGTGATGTTTCCCTTTGGGATACTGCTTTTTAACGTAAG
19AH AATCTAAGGCAATCGTCAAAAAGTGATGTTTCCCTTTGGGATACTGCTTTTTAACGTAAG
23FPO AATCTAAGGCAATCGTCAAAAAGTGATGTTTCCCTTTGGGATACTGCTTTTTAACGTAAG
19FTW AATCTAAGGCAATCGTCAAAAAGTGATGTTTCCCTTTGGGATACTGCTTTTTAACGTAAG
9VSP AATCTAAGGCAATCGTCAAAAAGTGATGTTTCCCTTTGGGATACTGCTTTTTAACGTAAG
TIGR4 AATCTAAGGCAATCGTCAAAAAGTGATGTTTCCCTTTGGGATACTGCTTTTTAACGTAAG
23FTW AATCTAAGGCAATCGTCAAAAAGTGATGTTTCCCTTTGGGATACTGCTTTTTAACGTAAG

14CSR GCAGGTATTCTTTTCGTTGTAATAATAATCAATGGCTCTGTCAAATGCTCCTCTGAAGGAG
670 GCAGGTATTCTTTTCGTTGTAATAATAATCAATGGCTCTGTCAAATGCTCCTCTGAAGGAG
6BF GCAGGTATTCTTTTCGTTGTAATAATAATCAATGGCTCTGTCAAATGCTCCTCTGAAGGAG
6BSP GCAGGTATTCTTTTCGTTGTAATAATAATCAATGGCTCTGTCAAATGCTCCTCTGAAGGAG
19AH GCAGGTATTCTTTTCGTTGTAATAATAATCAATGGCTCTGTCAAATGCTCCTCTGAAGGAG
23FPO GCAGGTATTCTTTTCGTTGTAATAATAATCAATGGCTCTGTCAAATGCTCCTCTGAAGGAG
19FTW GCAGGTATTCTTTTCGTTGTAATAATAATCAATGGCTCTGTCAAATGCTCCTCTGAAGGAG
9VSP GCAGGTATTCTTTTCGTTGTAATAATAATCAATGGCTCTGTCAAATGCTCCTCTGAAGGAG
TIGR4 GCAGGTATTCTTTTCGTTGTAATAATAATCAATGGCTCTGTCAAATGCTCCTCTGAAGGAG
23FTW GCAGGTATTCTTTTCGTTGTAATAATAATCAATGGCTCTGTCAAATGCTCCTCTGAAGGAG

14CSR GAGGACTAATTAGAATATTGTATCCTGTAACAGAGGCAACTTTGTGTCAGTAAAATCCGTA
670 GAGGACTAATTAGAATATTGTATCCTGTAACAGAGGCAACTTTGTGTCAGTAAAATCCGTA
6BF GAGGACTAATTAGAATATTGTATCCTGTAACAGAGGCAACTTTGTGTCAGTAAAATCCGTA
6BSP GAGGACTAATTAGAATATTGTATCCTGTAACAGAGGCAACTTTGTGTCAGTAAAATCCGTA
19AH GAGGACTAATTAGAATATTGTATCCTGTAACAGAGGCAACTTTGTGTCAGTAAAATCCGTA
23FPO GAGGACTAATTAGAATATTGTATCCTGTAACAGAGGCAACTTTGTGTCAGTAAAATCCGTA
19FTW GAGGACTAATTAGAATATTGTATCCTGTAACAGAGGCAACTTTGTGTCAGTAAAATCCGTA
9VSP GAGGACTAATTAGAATATTGTATCCTGTAACAGAGGCAACTTTGTGTCAGTAAAATCCGTA
TIGR4 GAGGACTAATTAGAATATTGTATCCTGTAACAGAGGCAACTTTGTGTCAGTAAAATCCGTA
23FTW GAGGACTAATTAGAATATTGTATCCTGTAACAGAGGCAACTTTGTGTCAGTAAAATCCGTA

14CSR AAATAATGGACTTTATTAAGTTTACATCTGCTTGATTATTTAAAATGATAAAAATCGGGA
670 AAATAATGGACTTTATTAAGTTTACATCTGCTTGATTATTTAAAATGATAAAAATCGGGA
6BF AAATAATGGACTTTATTAAGTTTACATCTGCTTGATTATTTAAAATGATAAAAATCGGGA
6BSP AAATAATGGACTTTATTAAGTTTACATCTGCTTGATTATTTAAAATGATAAAAATCGGGA
19AH AAATAATGGACTTTATTAAGTTTACATCTGCTTGATTATTTAAAATGATAAAAATCGGGA
23FPO AAATAATGGACTTTATTAAGTTTACATCTGCTTGATTATTTAAAATGATAAAAATCGGGA
19FTW AAATAATGGACTTTATTAAGTTTACATCTGCTTGATTATTTAAAATGATAAAAATCGGGA
9VSP AAATAATGGACTTTATTAAGTTTACATCTGCTTGATTATTTAAAATGATAAAAATCGGGA
TIGR4 AAATAATGGACTTTATTAAGTTTACATCTGCTTGATTATTTAAAATGATAAAAATCGGGA
23FTW AAATAATGGACTTTATTAAGTTTACATCTGCTTGATTATTTAAAATGATAAAAATCGGGA

Figure 196B

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14CSR TAGCAGGTAGTGAGGAAAAGATGGTTTCTGTCAAGTAGAGTGAGAAAAGGTACAGCCGAT
670 TAGCAGGTAGTGAGGAAAAGATGGTTTCTGTCAAGTAGAGTGAGAAAAGGTACAGCCGAT
6BF TAGCAGGTAGTGAGGAAAAGATGGTTTCTGTCAAGTAGAGTGAGAAAAGGTACAGCCGAT
6BSP TAGCAGGTAGTGAGGAAAAGATGGTTTCTGTCAAGTAGAGTGAGAAAAGGTACAGCCGAT
19AH TAGCAGGTAGTGAGGAAAAGATGGTTTCTGTCAAGTAGAGTGAGAAAAGGTACAGCCGAT
23FPO TAGCAGGTAGTGAGGAAAAGATGGTTTCTGTCAAGTAGAGTGAGAAAAGGTACAGCCGAT
19FTW TAGCAGGTAGTGAGGAAAAGATGGTTTCTGTCAAGTAGAGTGAGAAAAGGTACAGCCGAT
9VSP TAGCAGGTAGTGAGGAAAAGATGGTTTCTGTCAAGTAGAGTGAGAAAAGGTACAGCCGAT
TIGR4 TAGCAGGTAGTGAGGAAAAGATGGTTTCTGTCAAGTAGAGTGAGAAAAGGTACAGCCGAT
23FTW TAGCAGGTAGTGAGGAAAAGATGGTTTCTGTCAAGTAGAGTGAGAAAAGGTACAGCCGAT

14CSR GCTGGTCGATAACTCCTTCAATCTTCTGCTCAGTCATCCACTCTTGAACAATTGCTTTCG
670 GCTGGTCGATAACTCCTTCAATCTTCTGCTCAGTCATCCACTCTTGAACAATTGCTTTCG
6BF GCTGGTCGATAACTCCTTCAATCTTCTGCTCAGTCATCCACTCTTGAACAATTGCTTTCG
6BSP GCTGGTCGATAACTCCTTCAATCTTCTGCTCAGTCATCCACTCTTGAACAATTGCTTTCG
19AH GCTGGTCGATAACTCCTTCAATCTTCTGCTCAGTCATCCACTCTTGAACAATTGCTTTCG
23FPO GCTGGTCGATAACTCCTTCAATCTTCTGCTCAGTCATCCACTCTTGAACAATTGCTTTCG
19FTW GCTGGTCGATAACTCCTTCAATCTTCTGCTCAGTCATCCACTCTTGAACAATTGCTTTCG
9VSP GCTGGTCGATAACTCCTTCAATCTTCTGCTCAGTCATCCACTCTTGAACAATTGCTTTCG
TIGR4 GCTGGTCGATAACTCCTTCAATCTTCTGCTCAGTCATCCACTCTTGAACAATTGCTTTCG
23FTW GCTGGTCGATAACTCCTTCAATCTTCTGCTCAGTCATCCACTCTTGAACAATTGCTTTCG

14CSR AAATATGATACAGTGGCTTGTGCGCTTCAATCCCATAATGTTTCGTAATAATTATAATAGG
670 AAATATGATACAGTGGCTTGTGCGCTTCAATCCCATAATGTTTCGTAATAATTATAATAGG
6BF AAATATGATACAGTGGCTTGTGCGCTTCAATCCCATAATGTTTCGTAATAATTATAATAGG
6BSP AAATATGATACAGTGGCTTGTGCGCTTCAATCCCATAATGTTTCGTAATAATTATAATAGG
19AH AAATATGATACAGTGGCTTGTGCGCTTCAATCCCATAATGTTTCGTAATAATTATAATAGG
23FPO AAATATGATACAGTGGCTTGTGCGCTTCAATCCCATAATGTTTCGTAATAATTATAATAGG
19FTW AAATATGATACAGTGGCTTGTGCGCTTCAATCCCATAATGTTTCGTAATAATTATAATAGG
9VSP AAATATGATACAGTGGCTTGTGCGCTTCAATCCCATAATGTTTCGTAATAATTATAATAGG
TIGR4 AAATATGATACAGTGGCTTGTGCGCTTCAATCCCATAATGTTTCGTAATAATTATAATAGG
23FTW AAATATGATACAGTGGCTTGTGCGCTTCAATCCCATAATGTTTCGTAATAATTATAATAGG

14CSR GAACTAGATTTTGTAAACCAAACAAAACGTTCTTGTTAAGAAAGTCAGTGCTGTTAAAA
670 GAACTAGATTTTGTAAACCAAACAAAACGTTCTTGTTAAGAAAGTCAGTGCTGTTAAAA
6BF GAACTAGATTTTGTAAACCAAACAAAACGTTCTTGTTAAGAAAGTCAGTGCTGTTAAAA
6BSP GAACTAGATTTTGTAAACCAAACAAAACGTTCTTGTTAAGAAAGTCAGTGCTGTTAAAA
19AH GAACTAGATTTTGTAAACCAAACAAAACGTTCTTGTTAAGAAAGTCAGTGCTGTTAAAA
23FPO GAACTAGATTTTGTAAACCAAACAAAACGTTCTTGTTAAGAAAGTCAGTGCTGTTAAAA
19FTW GAACTAGATTTTGTAAACCAAACAAAACGTTCTTGTTAAGAAAGTCAGTGCTGTTAAAA
9VSP GAACTAGATTTTGTAAACCAAACAAAACGTTCTTGTTAAGAAAGTCAGTGCTGTTAAAA
TIGR4 GAACTAGATTTTGTAAACCAAACAAAACGTTCTTGTTAAGAAAGTCAGTGCTGTTAAAA
23FTW GAACTAGATTTTGTAAACCAAACAAAACGTTCTTGTTAAGAAAGTCAGTGCTGTTAAAA

14CSR AAGAAAGAGAATTTCGAAATGTCATTTCTTAAGATATTCTTGAACCTGGATAGTAGATGCT
670 AAGAAAGAGAATTTCGAAATGTCATTTCTTAAGATATTCTTGAACCTGGATAGTAGATGCT
6BF AAGAAAGAGAATTTCGAAATGTCATTTCTTAAGATATTCTTGAACCTGGATAGTAGATGCT
6BSP AAGAAAGAGAATTTCGAAATGTCATTTCTTAAGATATTCTTGAACCTGGATAGTAGATGCT
19AH AAGAAAGAGAATTTCGAAATGTCATTTCTTAAGATATTCTTGAACCTGGATAGTAGATGCT
23FPO AAGAAAGAGAATTTCGAAATGTCATTTCTTAAGATATTCTTGAACCTGGATAGTAGATGCT
19FTW AAGAAAGAGAATTTCGAAATGTCATTTCTTAAGATATTCTTGAACCTGGATAGTAGATGCT
9VSP AAGAAAGAGAATTTCGAAATGTCATTTCTTAAGATATTCTTGAACCTGGATAGTAGATGCT
TIGR4 AAGAAAGAGAATTTCGAAATGTCATTTCTTAAGATATTCTTGAACCTGGATAGTAGATGCT
23FTW AAGAAAGAGAATTTCGAAATGTCATTTCTTAAGATATTCTTGAACCTGGATAGTAGATGCT

Figure 196C

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14CSR      TTCCTCTTGTATGCTGAAGAATCAGTTGAATAGTATGAGTCTTTTTTTCTTGATTCCATT
670        TTCCTCTTGTATGCTGAAGAATCAGTTGAATAGTATGAGTCTTTTTTTCTTGATTCCATT
6BF        TTCCTCTTGTATGCTGAAGAATCAGTTGAATAGTATGAGTCTTTTTTTCTTGATTCCATT
6BSP       TTCCTCTTGTATGCTGAAGAATCAGTTGAATAGTATGAGTCTTTTTTTCTTGATTCCATT
19AH       TTCCTCTTGTATGCTGAAGAATCAGTTGAATAGTATGAGTCTTTTTTTCTTGATTCCATT
23FPO      TTCCTCTTGTATGCTGAAGAATCAGTTGAATAGTATGAGTCTTTTTTTCTTGATTCCATT
19FTW      TTCCTCTTGTATGCTGAAGAATCAGTTGAATAGTATGAGTCTTTTTTTCTTGATTCCATT
9VSP       TTCCTCTTGTATGCTGAAGAATCAGTTGAATAGTATGAGTCTTTTTTTCTTGATTCCATT
TIGR4      TTCCTCTTGTATGCTGAAGAATCAGTTGAATAGTATGAGTCTTTTTTTCTTGATTCCATT
23FTW      TTCCTCTTGTATGCTGAAGAATCAGTTGAATAGTATGAGTCTTTTTTTCTTGATTCCATT
*****

14CSR      TGTCCCTTGGAAAACGAAGAATTAGCAGAACAATAAACCAAAAAGATATAATCCAGTTCTT
670        TGTCCCTTGGAAAACGAAGAATTAGCAGAACAATAAACCAAAAAGATATAATCCAGTTCTT
6BF        TGTCCCTTGGAAAACGAAGAATTAGCAGAACAATAAACCAAAAAGATATAATCCAGTTCTT
6BSP       TGTCCCTTGGAAAACGAAGAATTAGCAGAACAATAAACCAAAAAGATATAATCCAGTTCTT
19AH       TGTCCCTTGGAAAACGAAGAATTAGCAGAACAATAAACCAAAAAGATATAATCCAGTTCTT
23FPO      TGTCCCTTGGAAAACGAAGAATTAGCAGAACAATAAACCAAAAAGATATAATCCAGTTCTT
19FTW      TGTCCCTTGGAAAACGAAGAATTAGCAGAACAATAAACCAAAAAGATATAATCCAGTTCTT
9VSP       TGTCCCTTGGAAAACGAAGAATTAGCAGAACAATAAACCAAAAAGATATAATCCAGTTCTT
TIGR4      TGTCCCTTGGAAAACGAAGAATTAGCAGAACAATAAACCAAAAAGATATAATCCAGTTCTT
23FTW      TGTCCCTTGGAAAACGAAGAATTAGCAGAACAATAAACCAAAAAGATATAATCCAGTTCTT
*****

14CSR      CCTGAGTAAAAGTCATGTTGGCATGTGGCTCTAAGTAAGTTTGGCAATGTTCCATCAAAA
670        CCTGAGTAAAAGTCATGTTGGCATGTGGCTCTAAGTAAGTTTGGCAATGTTCCATCAAAA
6BF        CCTGAGTAAAAGTCATGTTGGCATGTGGCTCTAAGTAAGTTTGGCAATGTTCCATCAAAA
6BSP       CCTGAGTAAAAGTCATGTTGGCATGTGGCTCTAAGTAAGTTTGGCAATGTTCCATCAAAA
19AH       CCTGAGTAAAAGTCATGTTGGCATGTGGCTCTAAGTAAGTTTGGCAATGTTCCATCAAAA
23FPO      CCTGAGTAAAAGTCATGTTGGCATGTGGCTCTAAGTAAGTTTGGCAATGTTCCATCAAAA
19FTW      CCTGAGTAAAAGTCATGTTGGCATGTGGCTCTAAGTAAGTTTGGCAATGTTCCATCAAAA
9VSP       CCTGAGTAAAAGTCATGTTGGCATGTGGCTCTAAGTAAGTTTGGCAATGTTCCATCAAAA
TIGR4      CCTGAGTAAAAGTCATGTTGGCATGTGGCTCTAAGTAAGTTTGGCAATGTTCCATCAAAA
23FTW      CCTGAGTAAAAGTCATGTTGGCATGTGGCTCTAAGTAAGTTTGGCAATGTTCCATCAAAA
*****

14CSR      TCGGATACATAAAGAGGTTTTTTAATTTTTCAAACCTCTTTGGACTCAGGGAACCTCAAGTG
670        TCGGATACATAAAGAGGTTTTTTAATTTTTCAAACCTCTTTGGACTCAGGGAACCTCAAGTG
6BF        TCGGATACATAAAGAGGTTTTTTAATTTTTCAAACCTCTTTGGACTCAGGGAACCTCAAGTG
6BSP       TCGGATACATAAAGAGGTTTTTTAATTTTTCAAACCTCTTTGGACTCAGGGAACCTCAAGTG
19AH       TCGGATACATAAAGAGGTTTTTTAATTTTTCAAACCTCTTTGGACTCAGGGAACCTCAAGTG
23FPO      TCGGATACATAAAGAGGTTTTTTAATTTTTCAAACCTCTTTGGACTCAGGGAACCTCAAGTG
19FTW      TCGGATACATAAAGAGGTTTTTTAATTTTTCAAACCTCTTTGGACTCAGGGAACCTCAAGTG
9VSP       TCGGATACATAAAGAGGTTTTTTAATTTTTCAAACCTCTTTGGACTCAGGGAACCTCAAGTG
TIGR4      TCGGATACATAAAGAGGTTTTTTAATTTTTCAAACCTCTTTGGACTCAGGGAACCTCAAGTG
23FTW      TCGGATACATAAAGAGGTTTTTTAATTTTTCAAACCTCTTTGGACTCAGGGAACCTCAAGTG
*****

14CSR      GAAATTCCCGACGTTTCCAAGTGAGTGCCACTAGTATGCTAAAATGAACATACTCGTCAG
670        GAAATTCCCGACGTTTCCAAGTGAGTGCCACTAGTATGCTAAAATGAACATACTCGTCAG
6BF        GAAATTCCCGACGTTTCCAAGTGAGTGCCACTAGTATGCTAAAATGAACATACTCGTCAG
6BSP       GAAATTCCCGACGTTTCCAAGTGAGTGCCACTAGTATGCTAAAATGAACATACTCGTCAG
19AH       GAAATTCCCGACGTTTCCAAGTGAGTGCCACTAGTATGCTAAAATGAACATACTCGTCAG
23FPO      GAAATTCCCGACGTTTCCAAGTGAGTGCCACTAGTATGCTAAAATGAACATACTCGTCAG
19FTW      GAAATTCCCGACGTTTCCAAGTGAGTGCCACTAGTATGCTAAAATGAACATACTCGTCAG
9VSP       GAAATTCCCGACGTTTCCAAGTGAGTGCCACTAGTATGCTAAAATGAACATACTCGTCAG
TIGR4      GAAATTCCCGACGTTTCCAAGTGAGTGCCACTAGTATGCTAAAATGAACATACTCGTCAG
23FTW      GAAATTCCCGACGTTTCCAAGTGAGTGCCACTAGTATGCTAAAATGAACATACTCGTCAG
*****

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14CSR GTGTGATTTCTAACAGTTCATGACTGAGTTGAGAATTAGACTGCACAATCATATGTGTGA
670 GTGTGATTTCTAACAGTTCATGACTGAGTTGAGAATTAGACTGCACAATCATATGTGTGA
6BF GTGTGATTTCTAACAGTTCATGACTGAGTTGAGAATTAGACTGCACAATCATATGTGTGA
6BSP GTGTGATTTCTAACAGTTCATGACTGAGTTGAGAATTAGACTGCACAATCATATGTGTGA
19AH GTGTGATTTCTAACAGTTCATGACTGAGTTGAGAATTAGACTGCACAATCATATGTGTGA
23FPO GTGTGATTTCTAACAGTTCATGACTGAGTTGAGAATTAGACTGCACAATCATATGTGTGA
19FTW GTGTGATTTCTAACAGTTCATGACTGAGTTGAGAATTAGACTGCACAATCATATGTGTGA
9VSP GTGTGATTTCTAACAGTTCATGACTGAGTTGAGAATTAGACTGCACAATCATATGTGTGA
TIGR4 GTGTGATTTCTAACAGTTCATGACTGAGTTGAGAATTAGACTGCACAATCATATGTGTGA
23FTW GTGTGATTTCTAACAGTTCATGACTGAGTTGAGAATTAGACTGCACAATCATATGTGTGA

14CSR CCCAATCCATACTTCCATCATTTCAAATCATAAATCTCAATACCAAAATGAAACTGGAGGA
670 CCCAATCCATACTTCCATCATTTCAAATCATAAATCTCAATACCAAAATGAAACTGGAGGA
6BF CCCAATCCATACTTCCATCATTTCAAATCATAAATCTCAATACCAAAATGAAACTGGAGGA
6BSP CCCAATCCATACTTCCATCATTTCAAATCATAAATCTCAATACCAAAATGAAACTGGAGGA
19AH CCCAATCCATACTTCCATCATTTCAAATCATAAATCTCAATACCAAAATGAAACTGGAGGA
23FPO CCCAATCCATACTTCCATCATTTCAAATCATAAATCTCAATACCAAAATGAAACTGGAGGA
19FTW CCCAATCCATACTTCCATCATTTCAAATCATAAATCTCAATACCAAAATGAAACTGGAGGA
9VSP CCCAATCCATACTTCCATCATTTCAAATCATAAATCTCAATACCAAAATGAAACTGGAGGA
TIGR4 CCCAATCCATACTTCCATCATTTCAAATCATAAATCTCAATACCAAAATGAAACTGGAGGA
23FTW CCCAATCCATACTTCCATCATTTCAAATCATAAATCTCAATACCAAAATGAAACTGGAGGA

14CSR GTGCAATTAATAAACGAATGCGATATT CAGGACCAACTACTTGATTTTTT CACAAGGTCCA
670 GTGCAATTAATAAACGAATGCGATATT CAGGACCAACTACTTGATTTTTT CACAAGGTCCA
6BF GTGCAATTAATAAACGAATGCGATATT CAGGACCAACTACTTGATTTTTT CACAAGGTCCA
6BSP GTGCAATTAATAAACGAATGCGATATT CAGGACCAACTACTTGATTTTTT CACAAGGTCCA
19AH GTGCAATTAATAAACGAATGCGATATT CAGGACCAACTACTTGATTTTTT CACAAGGTCCA
23FPO GTGCAATTAATAAACGAATGCGATATT CAGGACCAACTACTTGATTTTTT CACAAGGTCCA
19FTW GTGCAATTAATAAACGAATGCGATATT CAGGACCAACTACTTGATTTTTT CACAAGGTCCA
9VSP GTGCAATTAATAAACGAATGCGATATT CAGGACCAACTACTTGATTTTTT CACAAGGTCCA
TIGR4 GTGCAATTAATAAACGAATGCGATATT CAGGACCAACTACTTGATTTTTT CACAAGGTCCA
23FTW GTGCAATTAATAAACGAATGCGATATT CAGGACCAACTACTTGATTTTTT CACAAGGTCCA

14CSR AACCTACTGAACGTAGTAACAAGCCACACTTTTGTGTCGACGCGGTAGCCTGTTGCGATGG
670 AACCTACTGAACGTAGTAACAAGCCACACTTTTGTGTCGACGCGGTAGCCTGTTGCGATGG
6BF AACCTACTGAACGTAGTAACAAGCCACACTTTTGTGTCGACGCGGTAGCCTGTTGCGATGG
6BSP AACCTACTGAACGTAGTAACAAGCCACACTTTTGTGTCGACGCGGTAGCCTGTTGCGATGG
19AH AACCTACTGAACGTAGTAACAAGCCACACTTTTGTGTCGACGCGGTAGCCTGTTGCGATGG
23FPO AACCTACTGAACGTAGTAACAAGCCACACTTTTGTGTCGACGCGGTAGCCTGTTGCGATGG
19FTW AACCTACTGAACGTAGTAACAAGCCACACTTTTGTGTCGACGCGGTAGCCTGTTGCGATGG
9VSP AACCTACTGAACGTAGTAACAAGCCACACTTTTGTGTCGACGCGGTAGCCTGTTGCGATGG
TIGR4 AACCTACTGAACGTAGTAACAAGCCACACTTTTGTGTCGACGCGGTAGCCTGTTGCGATGG
23FTW AACCTACTGAACGTAGTAACAAGCCACACTTTTGTGTCGACGCGGTAGCCTGTTGCGATGG

14CSR AAATATACTCTTTTGTGTAAATTCGTTAAAGCTTTGATTACCTTGTAGTAGAAGAAGC
670 AAATATACTCTTTTGTGTAAATTCGTTAAAGCTTTGATTACCTTGTAGTAGAAGAAGC
6BF AAATATACTCTTTTGTGTAAATTCGTTAAAGCTTTGATTACCTTGTAGTAGAAGAAGC
6BSP AAATATACTCTTTTGTGTAAATTCGTTAAAGCTTTGATTACCTTGTAGTAGAAGAAGC
19AH AAATATACTCTTTTGTGTAAATTCGTTAAAGCTTTGATTACCTTGTAGTAGAAGAAGC
23FPO AAATATACTCTTTTGTGTAAATTCGTTAAAGCTTTGATTACCTTGTAGTAGAAGAAGC
19FTW AAATATACTCTTTTGTGTAAATTCGTTAAAGCTTTGATTACCTTGTAGTAGAAGAAGC
9VSP AAATATACTCTTTTGTGTAAATTCGTTAAAGCTTTGATTACCTTGTAGTAGAAGAAGC
TIGR4 AAATATACTCTTTTGTGTAAATTCGTTAAAGCTTTGATTACCTTGTAGTAGAAGAAGC
23FTW AAATATACTCTTTTGTGTAAATTCGTTAAAGCTTTGATTACCTTGTAGTAGAAGAAGC

Figure 196E

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14CSR GGAGTATTTTTAAAATAGTTGATTGGTTTATAAAGCTGATGGAAGTAATAATTCGTTTGAT
670 GGAGTATTTTTAAAATAGTTGATTGGTTTATAAAGCTGATGGAAGTAATAATTCGTTTGAT
6BF GGAGTATTTTTAAAATAGTTGATTGGTTTATAAAGCTGATGGAAGTAATAATTCGTTTGAT
6BSP GGAGTATTTTTAAAATAGTTGATTGGTTTATAAAGCTGATGGAAGTAATAATTCGTTTGAT
19AH GGAGTATTTTTAAAATAGTTGATTGGTTTATAAAGCTGATGGAAGTAATAATTCGTTTGAT
23FPO GGAGTATTTTTAAAATAGTTGATTGGTTTATAAAGCTGATGGAAGTAATAATTCGTTTGAT
19FTW GGAGTATTTTTAAAATAGTTGATTGGTTTATAAAGCTGATGGAAGTAATAATTCGTTTGAT
9VSP GGAGTATTTTTAAAATAGTTGATTGGTTTATAAAGCTGATGGAAGTAATAATTCGTTTGAT
TIGR4 GGAGTATTTTTAAAATAGTTGATTGGTTTATAAAGCTGATGGAAGTAATAATTCGTTTGAT
23FTW GGAGTATTTTTAAAATAGTTGATTGGTTTATAAAGCTGATGGAAGTAATAATTCGTTTGAT

14CSR GAGAATGGTGTTCGATTAAATGAACCTGTTGCGTATCTAAATTAAATGTCAACTCTTCCT
670 GAGAATGGTGTTCGATTAAATGAACCTGTTGCGTATCTAAATTAAATGTCAACTCTTCCT
6BF GAGAATGGTGTTCGATTAAATGAACCTGTTGCGTATCTAAATTAAATGTCAACTCTTCCT
6BSP GAGAATGGTGTTCGATTAAATGAACCTGTTGCGTATCTAAATTAAATGTCAACTCTTCCT
19AH GAGAATGGTGTTCGATTAAATGAACCTGTTGCGTATCTAAATTAAATGTCAACTCTTCCT
23FPO GAGAATGGTGTTCGATTAAATGAACCTGTTGCGTATCTAAATTAAATGTCAACTCTTCCT
19FTW GAGAATGGTGTTCGATTAAATGAACCTGTTGCGTATCTAAATTAAATGTCAACTCTTCCT
9VSP GAGAATGGTGTTCGATTAAATGAACCTGTTGCGTATCTAAATTAAATGTCAACTCTTCCT
TIGR4 GAGAATGGTGTTCGATTAAATGAACCTGTTGCGTATCTAAATTAAATGTCAACTCTTCCT
23FTW GAGAATGGTGTTCGATTAAATGAACCTGTTGCGTATCTAAATTAAATGTCAACTCTTCCT

14CSR CGAATGTTTCTTGTAATTCCTGCAAAATGCTTAGGAGACTTTTAGATTGTAATGAAGTTA
670 CGAATGTTTCTTGTAATTCCTGCAAAATGCTTAGGAGACTTTTAGATTGTAATGAAGTTA
6BF CGAATGTTTCTTGTAATTCCTGCAAAATGCTTAGGAGACTTTTAGATTGTAATGAAGTTA
6BSP CGAATGTTTCTTGTAATTCCTGCAAAATGCTTAGGAGACTTTTAGATTGTAATGAAGTTA
19AH CGAATGTTTCTTGTAATTCCTGCAAAATGCTTAGGAGACTTTTAGATTGTAATGAAGTTA
23FPO CGAATGTTTCTTGTAATTCCTGCAAAATGCTTAGGAGACTTTTAGATTGTAATGAAGTTA
19FTW CGAATGTTTCTTGTAATTCCTGCAAAATGCTTAGGAGACTTTTAGATTGTAATGAAGTTA
9VSP CGAATGTTTCTTGTAATTCCTGCAAAATGCTTAGGAGACTTTTAGATTGTAATGAAGTTA
TIGR4 CGAATGTTTCTTGTAATTCCTGCAAAATGCTTAGGAGACTTTTAGATTGTAATGAAGTTA
23FTW CGAATGTTTCTTGTAATTCCTGCAAAATGCTTAGGAGACTTTTAGATTGTAATGAAGTTA

14CSR AAGTAGACAGTTCATCTAGTTCAATAGACCGAATATCCAATAATATATTTAAAATGGTAA
670 AAGTAGACAGTTCATCTAGTTCAATAGACCGAATATCCAATAATATATTTAAAATGGTAA
6BF AAGTAGACAGTTCATCTAGTTCAATAGACCGAATATCCAATAATATATTTAAAATGGTAA
6BSP AAGTAGACAGTTCATCTAGTTCAATAGACCGAATATCCAATAATATATTTAAAATGGTAA
19AH AAGTAGACAGTTCATCTAGTTCAATAGACCGAATATCCAATAATATATTTAAAATGGTAA
23FPO AAGTAGACAGTTCATCTAGTTCAATAGACCGAATATCCAATAATATATTTAAAATGGTAA
19FTW AAGTAGACAGTTCATCTAGTTCAATAGACCGAATATCCAATAATATATTTAAAATGGTAA
9VSP AAGTAGACAGTTCATCTAGTTCAATAGACCGAATATCCAATAATATATTTAAAATGGTAA
TIGR4 AAGTAGACAGTTCATCTAGTTCAATAGACCGAATATCCAATAATATATTTAAAATGGTAA
23FTW AAGTAGACAGTTCATCTAGTTCAATAGACCGAATATCCAATAATATATTTAAAATGGTAA

14CSR TTTTATCTGTAATTCCTTTTTCAATGTATTTGTTTAGCATAGTTACCGAATCTTAGTTGC
670 TTTTATCTGTAATTCCTTTTTCAATGTATTTGTTTAGCATAGTTACCGAATCTTAGTTGC
6BF TTTTATCTGTAATTCCTTTTTCAATGTATTTGTTTAGCATAGTTACCGAATCTTAGTTGC
6BSP TTTTATCTGTAATTCCTTTTTCAATGTATTTGTTTAGCATAGTTACCGAATCTTAGTTGC
19AH TTTTATCTGTAATTCCTTTTTCAATGTATTTGTTTAGCATAGTTACCGAATCTTAGTTGC
23FPO TTTTATCTGTAATTCCTTTTTCAATGTATTTGTTTAGCATAGTTACCGAATCTTAGTTGC
19FTW TTTTATCTGTAATTCCTTTTTCAATGTATTTGTTTAGCATAGTTACCGAATCTTAGTTGC
9VSP TTTTATCTGTAATTCCTTTTTCAATGTATTTGTTTAGCATAGTTACCGAATCTTAGTTGC
TIGR4 TTTTATCTGTAATTCCTTTTTCAATGTATTTGTTTAGCATAGTTACCGAATCTTAGTTGC
23FTW TTTTATCTGTAATTCCTTTTTCAATGTATTTGTTTAGCATAGTTACCGAATCTTAGTTGC

Figure 196F

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14CSR ATATAGATAATTTTAATTATTATAATACAAAAGAACTAATTGTCTTGTCAAAAAGGTTG
670 ATATAGATAATTTTAATTATTATAATACAAAAGAACTAATTGTCTTGTCAAAAAGGTTG
6BF ATATAGATAATTTTAATTATTATAATACAAAAGAACTAATTGTCTTGTCAAAAAGGTTG
6BSP ATATAGATAATTTTAATTATTATAATACAAAAGAACTAATTGTCTTGTCAAAAAGGTTG
19AH ATATAGATAATTTTAATTATTATAATACAAAAGAACTAATTGTCTTGTCAAAAAGGTTG
23FPO ATATAGATAATTTTAATTATTATAATACAAAAGAACTAATTGTCTTGTCAAAAAGGTTG
19FTW ATATAGATAATTTTAATTATTATAATACAAAAGAACTAATTGTCTTGTCAAAAAGGTTG
9VSP ATATAGATAATTTTAATTATTATAATACAAAAGAACTAATTGTCTTGTCAAAAAGGTTG
TIGR4 ATATAGATAATTTTAATTATTATAATACAAAAGAACTAATTGTCTTGTCAAAAAGGTTG
23FTW ATATAGATAATTTTAATTATTATAATACAAAAGAACTAATTGTCTTGTCAAAAAGGTTG

14CSR TGGAATTTCCGACTTTATTGATAAAACAGCATGTAATAAAAGGCATTTTAAAGATAGTAA
670 TGGAATTTCCGACTTTATTGATAAAACAGCATGTAATAAAAGGCATTTTAAAGATAGTAA
6BF TGGAATTTCCGACTTTATTGATAAAACAGCATGTAATAAAAGGCATTTTAAAGATAGTAA
6BSP TGGAATTTCCGACTTTATTGATAAAACAGCATGTAATAAAAGGCATTTTAAAGATAGTAA
19AH TGGAATTTCCGACTTTATTGATAAAACAGCATGTAATAAAAGGCATTTTAAAGATAGTAA
23FPO TGGAATTTCCGACTTTATTGATAAAACAGCATGTAATAAAAGGCATTTTAAAGATAGTAA
19FTW TGGAATTTCCGACTTTATTGATAAAACAGCATGTAATAAAAGGCATTTTAAAGATAGTAA
9VSP TGGAATTTCCGACTTTATTGATAAAACAGCATGTAATAAAAGGCATTTTAAAGATAGTAA
TIGR4 TGGAATTTCCGACTTTATTGATAAAACAGCATGTAATAAAAGGCATTTTAAAGATAGTAA
23FTW TGGAATTTCCGACTTTATTGATAAAACAGCATGTAATAAAAGGCATTTTAAAGATAGTAA

14CSR TGAGTATTGGTGGAGTTTATGGCTTATTTTTTTTATTAGAAAATATTTTTTTATCAAAAT
670 TGAGTATTGGTGGAGTTTATGGCTTATTTTTTTTATTAGAAAATATTTTTTTATCAAAAT
6BF TGAGTATTGGTGGAGTTTATGGCTTATTTTTTTTATTAGAAAATATTTTTTTATCAAAAT
6BSP TGAGTATTGGTGGAGTTTATGGCTTATTTTTTTTATTAGAAAATATTTTTTTATCAAAAT
19AH TGAGTATTGGTGGAGTTTATGGCTTATTTTTTTTATTAGAAAATATTTTTTTATCAAAAT
23FPO TGAGTATTGGTGGAGTTTATGGCTTATTTTTTTTATTAGAAAATATTTTTTTATCAAAAT
19FTW TGAGTATTGGTGGAGTTTATGGCTTATTTTTTTTATTAGAAAATATTTTTTTATCAAAAT
9VSP TGAGTATTGGTGGAGTTTATGGCTTATTTTTTTTATTAGAAAATATTTTTTTATCAAAAT
TIGR4 TGAGTATTGGTGGAGTTTATGGCTTATTTTTTTTATTAGAAAATATTTTTTTATCAAAAT
23FTW TGAGTATTGGTGGAGTTTATGGCTTATTTTTTTTATTAGAAAATATTTTTTTATCAAAAT

14CSR ATTGTCGTTCTATAAAAAAATATGTGATAAAAAATATCTATTGTGATGGAAGTTGTTTTAA
670 ATTGTCGTTCTATAAAAAAATATGTGATAAAAAATATCTATTGTGATGGAAGTTGTTTTAA
6BF ATTGTCGTTCTATAAAAAAATATGTGATAAAAAATATCTATTGTGATGGAAGTTGTTTTAA
6BSP ATTGTCGTTCTATAAAAAAATATGTGATAAAAAATATCTATTGTGATGGAAGTTGTTTTAA
19AH ATTGTCGTTCTATAAAAAAATATGTGATAAAAAATATCTATTGTGATGGAAGTTGTTTTAA
23FPO ATTGTCGTTCTATAAAAAAATATGTGATAAAAAATATCTATTGTGATGGAAGTTGTTTTAA
19FTW ATTGTCGTTCTATAAAAAAATATGTGATAAAAAATATCTATTGTGATGGAAGTTGTTTTAA
9VSP ATTGTCGTTCTATAAAAAAATATGTGATAAAAAATATCTATTGTGATGGAAGTTGTTTTAA
TIGR4 ATTGTCGTTCTATAAAAAAATATGTGATAAAAAATATCTATTGTGATGGAAGTTGTTTTAA
23FTW ATTGTCGTTCTATAAAAAAATATGTGATAAAAAATATCTATTGTGATGGAAGTTGTTTTAA

14CSR TTTATACTAGGATAGTTAATAGTAATACTATACTATACTATATTGTATACAAGTGTGTCA
670 TTTATACTAGGATAGTTAATAGTAATACTATACTATACTATATTGTATACAAGTGTGTCA
6BF TTTATACTAGGATAGTTAATAGTAATACTATACTATACTATATTGTATACAAGTGTGTCA
6BSP TTTATACTAGGATAGTTAATAGTAATACTATACTATACTATATTGTATACAAGTGTGTCA
19AH TTTATACTAGGATAGTTAATAGTAATACTATACTATACTATATTGTATACAAGTGTGTCA
23FPO TTTATACTAGGATAGTTAATAGTAATACTATACTATACTATATTGTATACAAGTGTGTCA
19FTW TTTATACTAGGATAGTTAATAGTAATACTATACTATACTATATTGTATACAAGTGTGTCA
9VSP TTTATACTAGGATAGTTAATAGTAATACTATACTATACTATATTGTATACAAGTGTGTCA
TIGR4 TTTATACTAGGATAGTTAATAGTAATACTATACTATACTATATTGTATACAAGTGTGTCA
23FTW TTTATACTAGGATAGTTAATAGTAATACTATACTATACTA-----TATTGTATACAAGTGTGTCA

Figure 196G

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14CSR TTGCCAGGTTGAGAAGATAGCTATAACGCACCTTTTATACGCTTTTGCTACGTTTGTAGT
670 TTGCCAGGTTGAGAAGATAGCTATAACGCACCTTTTATACGCTTTTGCTACGTTTGTAGT
6BF TTGCCAGGTTGAGAAGATAGCTATAACGCACCTTTTATACGCTTTTGCTACGTTTGTAGT
6BSP TTGCCAGGTTGAGAAGATAGCTATAACGCACCTTTTATACGCTTTTGCTACGTTTGTAGT
19AH TTGCCAGGTTGAGAAGATAGCTATAACGCACCTTTTATACGCTTTTGCTACGTTTGTAGT
23FPO TTGCCAGGTTGAGAAGATAGCTATAACGCACCTTTTATACGCTTTTGCTACGTTTGTAGT
19FTW TTGCCAGGTTGAGAAGATAGCTATAACGCACCTTTTATACGCTTTTGCTACGTTTGTAGT
9VSP TTGCCAGGTTGAGAAGATAGCTATAACGCACCTTTTATACGCTTTTGCTACGTTTGTAGT
TIGR4 TTGCCAGGTTGAGAAGATAGCTATAACGCACCTTTTATACGCTTTTGCTACGTTTGTAGT
23FTW TTGCCAGGTTGAGAAGATAGCTATAACGCACCTTTTATACGCTTTTGCTACGTTTGTAGT

14CSR GAACGGATTAACCTCAG--TGAGATAAATTTTATCAGAACATAAGTAATCCGTTTCTTCGT
670 GAACGGATTAACCTCAG--TGAGATAAATTTTATCAGAACATAAGTAATCCGTTTCTTCGT
6BF GAACGGATTAACCTCAG--TGAGATAAATTTTATCAGAACATAAGTAATCCGTTTCTTCGT
6BSP GAACGGATTAACCTCAG--TGAGATAAATTTTATCAGAACATAAGTAATCCGTTTCTTCGT
19AH GAACGGATTAACCTCAGCATGAGATAAATTTTATCAGAA--TAAGTAATCCGTTTCTTCGT
23FPO GAACGGATTAACCTCAGCATGAGATAAATTTTATCAGAA--TAAGTAATCCGTTTCTTCGT
19FTW GAACGGATTAACCTCAG--TGAGATAAATTTTATCAGAACATAAGTAATCCGTTTCTTCGT
9VSP GAACGGATTAACCTCAG--TGAGATAAATTTTATCAGAACATAAGTAATCCGTTTCTTCGT
TIGR4 GAACGGATTAACCTCAG--TGAGATAAATTTTATCAGAACATAAGTAATCCGTTTCTTCGT
23FTW GAACGGATTAACCTCAG--TGAGATAAATTTTATCAGAACATAAGTAATCCGTTTCTTCGT

14CSR GTATACAGATTGAAAGTACCTATGAATCATAGAAGGATTAACCTGTTCTATGAATAATGC
670 GTATACAGATTGAAAGTACCTATGAATCATAGAAGGATTAACCTGTTCTATGAATAATGC
6BF GTATACAGATTGAAAGTACCTATGAATCATAGAAGGATTAACCTGTTCTATGAATAATGC
6BSP GTATACAGATTGAAAGTACCTATGAATCATAGAAGGATTAACCTGTTCTATGAATAATGC
19AH GTATACAGATTGAAAGTACCTATGAATCATAGAAGGATTAACCTGTTCTATGAATAATGC
23FPO GTATACAGATTGAAAGTACCTATGAATCATAGAAGGATTAACCTGTTCTATGAATAATGC
19FTW GTATACAGATTGAAAGTACCTATGAATCATAGAAGGATTAACCTGTTCTATGAATAATGC
9VSP GTATACAGATTGAAAGTACCTATGAATCATAGAAGGATTAACCTGTTCTATGAATAATGC
TIGR4 GTATACAGATTGAAAGTACCTATGAATCATAGAAGGATTAACCTGTTCTATGAATAATGC
23FTW GTATACAGATTGAAAGTACCTATGAATCATAGAAGGATTAACCTGTTCTATGAATAATGC

14CSR TTAACAGGGAGACACACATGAAAAAAGTAAGAAAGATATTTTTCAGAAAGGCAGTTGCAGGAC
670 TTAACAGGGAGACACACATGAAAAAAGTAAGAAAGATATTTTTCAGAAAGGCAGTTGCAGGAC
6BF TTAACAGGGAGACACACATGAAAAAAGTAAGAAAGATATTTTTCAGAAAGGCAGTTGCAGGAC
6BSP TTAACAGGGAGACACACATGAAAAAAGTAAGAAAGATATTTTTCAGAAAGGCAGTTGCAGGAC
19AH TTAACAGGGAGACACACATGAAAAAAGTAAGAAAGATATTTTTCAGAAAGGCAGTTGCAGGAC
23FPO TTAACAGGGAGACACACATGAAAAAAGTAAGAAAGATATTTTTCAGAAAGGCAGTTGCAGGAC
19FTW TTAACAGGGAGACACACATGAAAAAAGTAAGAAAGATATTTTTCAGAAAGGCAGTTGCAGGAC
9VSP TTAACAGGGAGACACACATGAAAAAAGTAAGAAAGATATTTTTCAGAAAGGCAGTTGCAGGAC
TIGR4 TTAACAGGGAGACACACATGAAAAAAGTAAGAAAGATATTTTTCAGAAAGGCAGTTGCAGGAC
23FTW TTAACAGGGAGACACACATGAAAAAAGTAAGAAAGATATTTTTCAGAAAGGCAGTTGCAGGAC

14CSR TGTGCTGTATATCTCAGTTGACAGCTTTTCTTCGATAGTTGCTTTAGCAGAAACGCCTG
670 TGTGCTGTATATCTCAGTTGACAGCTTTTCTTCGATAGTTGCTTTAGCAGAAACGCCTG
6BF TGTGCTGTATATCTCAGTTGACAGCTTTTCTTCGATAGTTGCTTTAGCAGAAACGCCTG
6BSP TGTGCTGTATATCTCAGTTGACAGCTTTTCTTCGATAGTTGCTTTAGCAGAAACGCCTG
19AH TGTGCTGTATATCTCAGTTGACAGCTTTTCTTCGATAGTTGCTTTAGCAGAAACGCCTG
23FPO TGTGCTGTATATCTCAGTTGACAGCTTTTCTTCGATAGTTGCTTTAGCAGAAACGCCTG
19FTW TGTGCTGTATATCTCAGTTGACAGCTTTTCTTCGATAGTTGCTTTAGCAGAAACGCCTG
9VSP TGTGCTGTATATCTCAGTTGACAGCTTTTCTTCGATAGTTGCTTTAGCAGAAACGCCTG
TIGR4 TGTGCTGTATATCTCAGTTGACAGCTTTTCTTCGATAGTTGCTTTAGCAGAAACGCCTG
23FTW TGTGCTGTATATCTCAGTTGACAGCTTTTCTTCGATAGTTGCTTTAGCAGAAACGCCTG

Figure 196H

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14CSR AAACCCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCGCTTC
670 AAACCCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCGCTTC
6BF AAACCCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCGCTTC
6BSP AAACCCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCGCTTC
19AH AAACCCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCGCTTC
23FPO AAACCCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCGCTTC
19FTW AAACCCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCGCTTC
9VSP AAACCCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCGCTTC
TIGR4 AAACCCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCGCTTC
23FTW AAACCCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCGCTTC

14CSR TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGGATGGCACAACCTGTTTCGCAAAGGA
670 TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGGATGGCACAACCTGTTTCGCAAAGGA
6BF TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGGATGGCACAACCTGTTTCGCAAAGGA
6BSP TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGGATGGCACAACCTGTTTCGCAAAGGA
19AH TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGGATGGCACAACCTGTTTCGCAAAGGA
23FPO TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGGATGGCACAACCTGTTTCGCAAAGGA
19FTW TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGGATGGCACAACCTGTTTCGCAAAGGA
9VSP TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGAATGGCACAACCTGTTTCGCAAAGGA
TIGR4 TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGGATGGCACAACCTGTTTCGCAAAGGA
23FTW TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGGATGGCACAACCTGTTTCGCAAAGGA

14CSR CAGAGGCGCAAAACAGGAGAAGCGATATTTTCAAACATAAAACCTGGGACATACACCTTGA
670 CAGAGGCGCAAAACAGGAGAAGCGATATTTTCAAACATAAAACCTGGGACATACACCTTGA
6BF CAGAGGCGCAAAACAGGAGAAGCGATATTTTCAAACATAAAACCTGGGACATACACCTTGA
6BSP CAGAGGCGCAAAACAGGAGAAGCGATATTTTCAAACATAAAACCTGGGACATACACCTTGA
19AH CAGAGGCGCAAAACAGGAGAAGCGATATTTTCAAACATAAAACCTGGGACATACACCTTGA
23FPO CAGAGGCGCAAAACAGGAGAAGCGATATTTTCAAACATAAAACCTGGGACATACACCTTGA
19FTW CAGAGGCGCAAAACAGGAGAAGCGATATTTTCAAACATAAAACCTGGGACATACACCTTGA
9VSP CAGAGGCGCAAAACAGGAGAAGCGATATTTTCAAACATAAAACCTGGGACATACACCTTGA
TIGR4 CAGAGGCGCAAAACAGGAGAAGCGATATTTTCAAACATAAAACCTGGGACATACACCTTGA
23FTW CAGAGGCGCAAAACAGGAGAAGCGATATTTTCAAACATAAAACCTGGGACATACACCTTGA

14CSR CAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACAATGGACTGTTGAAGTTG
670 CAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACAATGGACTGTTGAAGTTG
6BF CAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACAATGGACTGTTGAAGTTG
6BSP CAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACAATGGACTGTTGAAGTTG
19AH CAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACAATGGACTGTTGAAGTTG
23FPO CAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACAATGGACTGTTGAAGTTG
19FTW CAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACAATGGACTGTTGAAGTTG
9VSP CAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACAACGGACTGTTGAAGTTG
TIGR4 CAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACAATGGACTGTTGAAGTTG
23FTW CAGAAGCCCAACCTCCAGTTGGTTATAAACCTCTACTAAACAATGGACTGTTGAAGTTG

14CSR AGAAGAATGGTTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTAT
670 AGAAGAATGGTTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTAT
6BF AGAAGAATGGTTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTAT
6BSP AGAAGAATGGTTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTAT
19AH AGAAGAATGGTTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTAT
23FPO AGAAGAATGGTTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTAT
19FTW AGAAGAATGGTTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTAT
9VSP AGAAGAATGGTTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTAT
TIGR4 AGAAGAATGGTTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTAT
23FTW AGAAGAATGGTTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTAT

14CSR CTGACCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTA
670 CTGACCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTA
6BF CTGACCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTA
6BSP CTGACCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTA
19AH CTGACCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTA
23FPO CTGACCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTA
19FTW CTGACCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTA
9VSP CTGACCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTA
TIGR4 CTGACCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTA
23FTW CTGACCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTA

14CSR AGGTAGATGGTTCGGAAAAAACGGACAGCACAAGGCGTTGAATCCGAATCCATATGAAC
670 AGGTAGATGGTTCGGAAAAAACGGACAGCACAAGGCGTTGAATCCGAATCCATATGAAC
6BF AGGTAGATGGTTCGGAAAAAACGGACAGCACAAGGCGTTGAATCCGAATCCATATGAAC
6BSP AGGTAGATGGTTCGGAAAAAACGGACAGCACAAGGCGTTGAATCCGAATCCATATGAAC
19AH AGGTAGATGGTTCGGAAAAAACGGACAGCACAAGGCGTTGAATCCGAATCCATATGAAC
23FPO AGGTAGATGGTTCGGAAAAAACGGACAGCACAAGGCGTTGAATCCGAATCCATATGAAC
19FTW AGGTAGATGGTTCGGAAAAAACGGACAGCACAAGGCGTTGAATCCGAATCCATATGAAC
9VSP AGGTAGATGGTTCGGAAAAAACGGACAGCACAAGGCGTTGAATCCGAATCCATATGAAC
TIGR4 AGGTAGATGGTTCGGAAAAAACGGACAGCACAAGGCGTTGAATCCGAATCCATATGAAC
23FTW AGGTAGATGGTTCGGAAAAAACGGACAGCACAAGGCGTTGAATCCGAATCCATATGAAC

14CSR GTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATAATTTGGATGATA
670 GTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATAATTTGGATGATA
6BF GTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATAATTTGGATGATA
6BSP GTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATAATTTGGATGATA
19AH GTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATAATTTGGATGATA
23FPO GTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATAATTTGGATGATA
19FTW GTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATAATTTGGATGATA
9VSP GTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATAATTTGGATGATA
TIGR4 GTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATAATTTGGATGATA
23FTW GTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATAATTTGGATGATA

14CSR ACCAATATGGAATCGAGTTGACGGTTAGTGGTAAAACGACGGTTGAAACGAAAGAAGCCT
670 ACCAATATGGAATCGAGTTGACGGTTAGTGGTAAAACGACGGTTGAAACGAAAGAAGCCT
6BF ACCAATATGGAATCGAGTTGACGGTTAGTGGTAAAACGACGGTTGAAACGAAAGAAGCCT
6BSP ACCAATATGGAATCGAGTTGACGGTTAGTGGTAAAACGACGGTTGAAACGAAAGAAGCCT
19AH ACCAATATGGAATCGAGTTGACGGTTAGTGGTAAAACGACGGTTGAAACGAAAGAAGCCT
23FPO ACCAATATGGAATCGAATTGACGGTTAGTGGGAAAACAGTGTATGAACGAAAAGATAAGT
19FTW ACCAATATGGAATCGAATTGACGGTTAGTGGGAAAACAGTGTATGAACGAAAAGATAAGT
9VSP ACCAATATGGAATCGAATTGACGGTTAGTGGGAAAACAGTGTATGAACGAAAAGATAAGT
TIGR4 ACCAATATGGAATCGAATTGACGGTTAGTGGGAAAACAGTGTATGAACGAAAAGATAAGT
23FTW ACCAATATGGAATCGAATTGACGGTTAGTGGGAAAACAGTGTATGAACGAAAAGATAAGT
***** * **** *

14CSR CTACTCCGCTAGATGTTGTTATTCTATTAGATAACTCCAATAGTATGAGTAATATTCGAC
670 CTACTCCGCTAGATGTTGTTATTCTATTAGATAACTCCAATAGTATGAGTAATATTCGAC
6BF CTACTCCGCTAGATGTTGTTATTCTATTAGATAACTCCAATAGTATGAGTAATATTCGAC
6BSP CTACTCCGCTAGATGTTGTTATTCTATTAGATAACTCCAATAGTATGAGTAATATTCGAC
19AH CTACTCCGCTAGATGTTGTTATTCTATTAGATAACTCCAATAGTATGAGTAATATTCGAC
23FPO CTACTCCGCTAGATGTTGTTATTCTATTAGATAACTCCAATAGTATGAGTAATATTCGAC
19FTW CTGTGCCGCTGGATGTCGTTATCTTGCTCGATAACTCAAATAGTATGAGTAACATTTCGAA
9VSP CTGTGCCGCTGGATGTCGTTATCTTGCTCGATAACTCAAATAGTATGAGTAACATTTCGAA
TIGR4 CTGTGCCGCTGGATGTCGTTATCTTGCTCGATAACTCAAATAGTATGAGTAACATTTCGAA
23FTW CTGTGCCGCTGGATGTCGTTATCTTGCTCGATAACTCAAATAGTATGAGTAACATTTCGAA
** ***** *

Figure 196J

Figure 196K

PCT/US05/27239 404/487

14CSR CAGAGGAATTGAACAAAGACAAATTGATGTATCAATTCGGCGCGACTTTTACCCAGAAGG
670 CAGAGGAATTGAACAAAGACAAATTGATGTATCAATTCGGCGCGACTTTTACCCAGAAGG
6BF CAGAGGAATTGAACAAAGACAAATTGATGTATCAATTCGGCGCGACTTTTACCCAGAAGG
6BSP CAGAGGAATTGAACAAAGACAAATTGATGTATCAATTCGGCGCGACTTTTACCCAGAAGG
19AH CAGAGGAATTGAACAAAGACAAATTGATGTATCAATTCGGCGCGACTTTTACCCAGAAGG
23FPO CAGAGGAATTGAACAAAGACAAATTGATGTATCAATTCGGCGCGACTTTTACCCAGAAGG
19FTW CAGAAGATCATGATGGAATAGATTGATGTACCAATTCGGTGCCACTTTTACTCAGAAAG
9VSP CAGAAGACCATGATGGAATAGATTGATGTACCAATTCGGTGCCACTTTTACTCAGAAAG
TIGR4 CAGAAGACCATGATGGAATAGATTGATGTACCAATTCGGTGCCACTTTTACTCAGAAAG
23FTW CAGAAGACCATGATGGAATAGATTGATGTACCAATTCGGTGCCACTTTTACTCAGAAAG
***** * * * *
14CSR CTTTGATGACCGCTGATGATATCTTGACAAAGCAGGCAAGACCAAACAGTAAAAAGGTTA
670 CTTTGATGACCGCTGATGATATCTTGACAAAGCAGGCAAGACCAAACAGTAAAAAGGTTA
6BF CTTTGATGACCGCTGATGATATCTTGACAAAGCAGGCAAGACCAAACAGTAAAAAGGTTA
6BSP CTTTGATGACCGCTGATGATATCTTGACAAAGCAGGCAAGACCAAACAGTAAAAAGGTTA
19AH CTTTGATGACCGCTGATGATATCTTGACAAAGCAGGCAAGACCAAACAGTAAAAAGGTTA
23FPO CTTTGATGACCGCTGATGATATCTTGACAAAGCAGGCAAGACCAAACAGTAAAAAGGTTA
19FTW CTTTGATGAAGGCAGATGAGATTTTGACACAACAAGCGAGACAAAATAGTCAAAAAGTCA
9VSP CTTTGATGAAGGCCGATGAGATTTTGACACAACAAGCGAGACAAAATAGTCAAAAAGTCA
TIGR4 CTTTGATGAAGGCAGATGAGATTTTGACACAACAAGCGAGACAAAATAGTCAAAAAGTCA
23FTW CTTTGATGAAGGCAGATGAGATTTTGACACAACAAGCGAGACAAAATAGTCAAAAAGTCA
***** * * * *
14CSR TTTTCCACATTACAGATGGTGTTCGACTATGTCATATCCAATTAATTTTAAATATACAG
670 TTTTCCACATTACAGATGGTGTTCGACTATGTCATATCCAATTAATTTTAAATATACAG
6BF TTTTCCACATTACAGATGGTGTTCGACTATGTCATATCCAATTAATTTTAAATATACAG
6BSP TTTTCCACATTACAGATGGTGTTCGACTATGTCATATCCAATTAATTTTAAATATACAG
19AH TTTTCCACATTACAGATGGTGTTCGACTATGTCATATCCAATTAATTTTAAATATACAG
23FPO TTTTCCACATTACAGATGGTGTTCGACTATGTCATATCCAATTAATTTTAAATATACAG
19FTW TTTTCCATATTACGGATGGTGTCCCAACTATGTCGATCCGATTAATTTTAAATCATGCTA
9VSP TTTTCCATATTACGGATGGTGTCCCAACTATGTCGATCCGATTAATTTTAAATCATGCTA
TIGR4 TTTTCCATATTACGGATGGTGTCCCAACTATGTCGATCCGATTAATTTTAAATCATGCTA
23FTW TTTTCCATATTACGGATGGTGTCCCAACTATGTCGATCCGATTAATTTTAAATCATGCTA
***** * * * *
14CSR GAACGACGCAATCGTACAGAACTCAGCTGAATA-ATTTTAAAGCAAAAACCTCCAAATAGT
670 GAACGACGCAATCGTACAGAACTCAGCTGAATA-ATTTTAAAGCAAAAACCTCCAAATAGT
6BF GAACGACGCAATCGTACAGAACTCAGCTGAATA-ATTTTAAAGCAAAAACCTCCAAATAGT
6BSP GAACGACGCAATCGTACAGAACTCAGCTGAATA-ATTTTAAAGCAAAAACCTCCAAATAGT
19AH GAACGACGCAATCGTACAGAACTCAGCTGAATA-ATTTTAAAGCAAAAACCTCCAAATAGT
23FPO GAACGACGCAATCGTACAGAACTCAGCTGAATA-ATTTTAAAGCAAAAACCTCCAAATAGT
19FTW CGTTTGCTCCATCATATCAAAATCAACTAAATGCATTTTTTAGTAAAT-CTCCTAATAAA
9VSP CGTTTGCTCCATCATATCAAAATCAACTAAATGCATTTTTTAGTAAAT-CTCCTAATAAA
TIGR4 CGTTTGCTCCATCATATCAAAATCAACTAAATGCATTTTTTAGTAAAT-CTCCTAATAAA
23FTW CGTTTGCTCCATCATATCAAAATCAACTAAATGCATTTTTTAGTAAAT-CTCCTAATAAA
* * * * *
14CSR AGCGGGATATTACTGGAGGACTTTGTTACATGGTCAGCAGATGGTGAACATAAGATTGTT
670 AGCGGGATATTACTGGAGGACTTTGTTACATGGTCAGCAGATGGTGAACATAAGATTGTT
6BF AGCGGGATATTACTGGAGGACTTTGTTACATGGTCAGCAGATGGTGAACATAAGATTGTT
6BSP AGCGGGATATTACTGGAGGACTTTGTTACATGGTCAGCAGATGGTGAACATAAGATTGTT
19AH AGCGGGATATTACTGGAGGACTTTGTTACATGGTCAGCAGATGGTGAACATAAGATTGTT
23FPO AGCGGGATATTACTGGAGGACTTTGTTACATGGTCAGCAGATGGTGAACATAAGATTGTT
19FTW GATGGAATACTATTAAAGTGATTTTATTACGCAAGCAACTAGTGGAGAACATACAATTGTA
9VSP GATGGAATACTATTAAAGTGATTTTATTACGCAAGCAACTAGTGGAGAACATACAATTGTA
TIGR4 GATGGAATACTATTAAAGTGATTTTATTACGCAAGCAACTAGTGGAGAACATACAATTGTA
23FTW GATGGAATACTATTAAAGTGATTTTATTACGCAAGCAACTAGTGGAGAACATACAATTGTA
* * * * *

Figure 196L

14CSR CGTGGAGATGGTGAAAGTTATCAGATGTTTACGAAGAAACCTGT-----AACAGACCAA
670 CGTGGAGATGGTGAAAGTTATCAGATGTTTACGAAGAAACCTGT-----AACAGACCAA
6BF CGTGGAGATGGTGAAAGTTATCAGATGTTTACGAAGAAACCTGT-----AACAGACCAA
6BSP CGTGGAGATGGTGAAAGTTATCAGATGTTTACGAAGAAACCTGT-----AACAGACCAA
19AH CGTGGAGATGGTGAAAGTTATCAGATGTTTACGAAGAAACCTGT-----AACAGACCAA
23FPO CGTGGAGATGGTGAAAGTTATCAGATGTTTACGAAGAAACCTGT-----AACAGACCAA
19FTW CGCGGAGATGGGCAAAAGTTACCAGATGTTTACAGATAAGACAGTTTTATGAAAAAGGTGCT
9VSP CGCGGAGATGGGCAAAAGTTACCAGATGTTTACAGATAAGACAGTTTTATGAAAAAGGTGCT
TIGR4 CGCGGAGATGGGCAAAAGTTACCAGATGTTTACAGATAAGACAGTTTTATGAAAAAGGTGCT
23FTW CGCGGAGATGGGCAAAAGTTACCAGATGTTTACAGATAAGACAGTTTTATGAAAAAGGTGCT
** * ** * ** * ** * ** * ** *

14CSR TACGGAGTTCATCAAAAT---ACTTTCAATCACCTCCATGGAGCAGAGAGCTAAATTAGTT
670 TACGGAGTTCATCAAAAT---ACTTTCAATCACCTCCATGGAGCAGAGAGCTAAATTAGTT
6BF TACGGAGTTCATCAAAAT---ACTTTCAATCACCTCCATGGAGCAGAGAGCTAAATTAGTT
6BSP TACGGAGTTCATCAAAAT---ACTTTCAATCACCTCCATGGAGCAGAGAGCTAAATTAGTT
19AH TACGGAGTTCATCAAAAT---ACTTTCAATCACCTCCATGGAGCAGAGAGCTAAATTAGTT
23FPO TACGGAGTTCATCAAAAT---ACTTTCAATCACCTCCATGGAGCAGAGAGCTAAATTAGTT
19FTW CCTGCGAGCTTTCCCAGTTAAACCTGAAAAATATTCTGAAATGAAGCGGCTGGTTTATGCA
9VSP CCTGCGAGCTTTCCCAGTTAAACCTGAAAAATATTCTGAAATGAAGCGGCTGGTTTATGCA
TIGR4 CCTGCGAGCTTTCCCAGTTAAACCTGAAAAATATTCTGAAATGAAGCGGCTGGTTTATGCA
23FTW CCTGCGAGCTTTCCCAGTTAAACCTGAAAAATATTCTGAAATGAAGCGGCTGGTTTATGCA
* * * * * * * * * * * * * * *

14CSR TCAGCGGGATATAGGTTCTATGGAACCTGACTTGTATTTATATTGGCGTGATAGTATTCTA
670 TCAGCGGGATATAGGTTCTATGGAACCTGACTTGTATTTATATTGGCGTGATAGTATTCTA
6BF TCAGCGGGATATAGGTTCTATGGAACCTGACTTGTATTTATATTGGCGTGATAGTATTCTA
6BSP TCAGCGGGATATAGGTTCTATGGAACCTGACTTGTATTTATATTGGCGTGATAGTATTCTA
19AH TCAGCGGGATATAGGTTCTATGGAACCTGACTTGTATTTATATTGGCGTGATAGTATTCTA
23FPO TCAGCGGGATATAGGTTCTATGGAACCTGACTTGTATTTATATTGGCGTGATAGTATTCTA
19FTW GTTTATAGGCGATCCAATTAATGGTGGATATATTGGCTTAATTGGAGAGAGAGTATTCTG
9VSP GTTTATAGGCGATCCAATTAATGGTGGATATATTGGCTTAATTGGAGAGAGAGTATTCTG
TIGR4 GTTTATAGGCGATCCAATTAATGGTGGATATATTGGCTTAATTGGAGAGAGAGTATTCTG
23FTW GTTTATAGGCGATCCAATTAATGGTGGATATATTGGCTTAATTGGAGAGAGAGTATTCTG
* * * * * * * * * * * * * * *

14CSR GCCTATCCATTTAACTCTAGTAGCCGATTGGATTACCAACCATGGTGACCCTACGACTTGG
670 GCCTATCCATTTAACTCTAGTAGCCGATTGGATTACCAACCATGGTGACCCTACGACTTGG
6BF GCCTATCCATTTAACTCTAGTAGCCGATTGGATTACCAACCATGGTGACCCTACGACTTGG
6BSP GCCTATCCATTTAACTCTAGTAGCCGATTGGATTACCAACCATGGTGACCCTACGACTTGG
19AH GCCTATCCATTTAACTCTAGTAGCCGATTGGATTACCAACCATGGTGACCCTACGACTTGG
23FPO GCCTATCCATTTAACTCTAGTAGCCGATTGGATTACCAACCATGGTGACCCTACGACTTGG
19FTW GCTTATCCGTTTAACTCTAATACTGCTAAAATTACCAATCATGGTGACCCTACAAGATGG
9VSP GCTTATCCGTTTAACTCTAATACTGCTAAAATTACCAATCATGGTGACCCTACAAGATGG
TIGR4 GCTTATCCGTTTAACTCTAATACTGCTAAAATTACCAATCATGGTGACCCTACAAGATGG
23FTW GCTTATCCGTTTAACTCTAATACTGCTAAAATTACCAATCATGGTGACCCTACAAGATGG
* * * * * * * * * * * * * * *

14CSR TATTATAACGGAAATATGGCTCAGGATGGCTATGATGTCTTCACTGTTGGGGTTGGTGTA
670 TATTATAACGGAAATATGGCTCAGGATGGCTATGATGTCTTCACTGTTGGGGTTGGTGTA
6BF TATTATAACGGAAATATGGCTCAGGATGGCTATGATGTCTTCACTGTTGGGGTTGGTGTA
6BSP TATTATAACGGAAATATGGCTCAGGATGGCTATGATGTCTTCACTGTTGGGGTTGGTGTA
19AH TATTATAACGGAAATATGGCTCAGGATGGCTATGATGTCTTCACTGTTGGGGTTGGTGTA
23FPO TATTATAACGGAAATATGGCTCAGGATGGCTATGATGTCTTCACTGTTGGGGTTGGTGTA
19FTW TACTATAACGGGAATATTGCTCCTGATGGGTATGATGTCTTTACGGTAGGTATTGGTATT
9VSP TACTATAACGGGAATATTGCTCCTGATGGGTATGATGTCTTTACGGTAGGTATTGGTATT
TIGR4 TACTATAACGGGAATATTGCTCCTGATGGGTATGATGTCTTTACGGTAGGTATTGGTATT
23FTW TACTATAACGGGAATATTGCTCCTGATGGGTATGATGTCTTTACGGTAGGTATTGGTATT
* * * * * * * * * * * * * * *

Figure 196M

14CSR AACCAGGGGATCCTGGTACGGATGAAGCAACCGGCTACTAGATTTATGCAGAGCATCTCTAGT
670 AACCGGGGATCCTGGTACGGATGAAGCAACCGGCTACTAGATTTATGCAGAGCATCTCTAGT
6BF AACCGGGGATCCTGGTACGGATGAAGCAACCGGCTACTAGATTTATGCAGAGCATCTCTAGT
6BSP AACCGGGGATCCTGGTACGGATGAAGCAACCGGCTACTAGATTTATGCAGAGCATCTCTAGT
19AH AACCGGGGATCCTGGTACGGATGAAGCAACCGGCTACTAGATTTATGCAGAGCATCTCTAGT
23FPO AACCGGGGATCCTGGTACGGATGAAGCAACCGGCTACTAGATTTATGCAGAGCATCTCTAGT
19FTW AACCGGAGATCCTGGTACGGATGAAGCAACCGGCTACTAGTTTTATGCAAAGTATTTCTAGT
9VSP AACCGGAGATCCTGGTACGGATGAAGCAACCGGCTACTAGTTTTATGCAAAGTATTTCTAGT
TIGR4 AACCGGAGATCCTGGTACGGATGAAGCAACCGGCTACTAGTTTTATGCAAAGTATTTCTAGT
23FTW AACCGGAGATCCTGGTACGGATGAAGCAACCGGCTACTAGTTTTATGCAAAGTATTTCTAGT

14CSR TCTCCTGACAACCTACACTAACGTAGCAGATCCATCTCAGATTTTACAAGAATTGAATCGC
670 TCTCCTGACAACCTACACTAACGTAGCAGATCCATCTCAGATTTTACAAGAATTGAATCGC
6BF TCTCCTGACAACCTACACTAACGTAGCAGATCCATCTCAGATTTTACAAGAATTGAATCGC
6BSP TCTCCTGACAACCTACACTAACGTAGCAGATCCATCTCAGATTTTACAAGAATTGAATCGC
19AH TCTCCTGACAACCTACACTAACGTAGCAGATCCATCTCAGATTTTACAAGAATTGAATCGC
23FPO TCTCCTGACAACCTACACTAACGTAGCAGATCCATCTCAGATTTTACAAGAATTGAATCGC
19FTW AAACCTGAAAACCTATACCAATGTTACTGACACGACAAAAATATTGGAACAGTTGAATCGT
9VSP AAACCTGAAAACCTATACCAATGTTACTGACACGACAAAAATATTGGAACAGTTGAATCGT
TIGR4 AAACCTGAAAACCTATACCAATGTTACTGACACGACAAAAATATTGGAACAGTTGAATCGT
23FTW AAACCTGAAAACCTATACCAATGTTACTGACACGACAAAAATATTGGAACAGTTGAATCGT

14CSR TACTTCTATACTATCGTCAATGAGAAGAAATCTATCGAAAATGGTACGATTACAGACCCG
670 TACTTCTATACTATCGTCAATGAGAAGAAATCTATCGAAAATGGTACGATTACAGACCCG
6BF TACTTCTATACTATCGTCAATGAGAAGAAATCTATCGAAAATGGTACGATTACAGACCCG
6BSP TACTTCTATACTATCGTCAATGAGAAGAAATCTATCGAAAATGGTACGATTACAGACCCG
19AH TACTTCTATACTATCGTCAATGAGAAGAAATCTATCGAAAATGGTACGATTACAGACCCG
23FPO TACTTCTATACTATCGTCAATGAGAAGAAATCTATCGAAAATGGTACGATTACAGACCCG
19FTW TATTTCCACACCATCGTAACTGAAAAGAAATCAATTGAGAATGGTACGATTACAGATCCG
9VSP TATTTCCACACCATCGTAACTGAAAAGAAATCAATTGAGAATGGTACGATTACAGATCCG
TIGR4 TATTTCCACACCATCGTAACTGAAAAGAAATCAATTGAGAATGGTACGATTACAGATCCG
23FTW TATTTCCACACCATCGTAACTGAAAAGAAATCAATTGAGAATGGTACGATTACAGATCCG
** * * * * *

14CSR ATGGGTGAACTAATTGATTTCCAATTGGGAGCAGATGGAAGGTTTGATCCAGCGGATTAC
670 ATGGGTGAACTAATTGATTTCCAATTGGGAGCAGATGGAAGGTTTGATCCAGCGGATTAC
6BF ATGGGTGAACTAATTGATTTCCAATTGGGAGCAGATGGAAGGTTTGATCCAGCGGATTAC
6BSP ATGGGTGAACTAATTGATTTCCAATTGGGAGCAGATGGAAGGTTTGATCCAGCGGATTAC
19AH ATGGGTGAACTAATTGATTTCCAATTGGGAGCAGATGGAAGGTTTGATCCAGCGGATTAC
23FPO ATGGGTGAACTAATTGATTTCCAATTGGGAGCAGATGGAAGGTTTGATCCAGCGGATTAC
19FTW ATGGGTGAGTTAATTGATTTGCAATTGGGCACAGATGGAAGATTGATCCAGCAGATTAC
9VSP ATGGGTGAGTTAATTGATTTGCAATTGGGCACAGATGGAAGATTGATCCAGCAGATTAC
TIGR4 ATGGGTGAGTTAATTGATTTGCAATTGGGCACAGATGGAAGATTGATCCAGCAGATTAC
23FTW ATGGGTGAGTTAATTGATTTGCAATTGGGCACAGATGGAAGATTGATCCAGCAGATTAC

14CSR ACTTTAACTGCAAACGATGGTAGTTCGTTGGTGAATAATGTCCCTACTGGGGGACCACAA
670 ACTTTAACTGCAAACGATGGTAGTTCGTTGGTGAATAATGTCCCTACTGGGGGACCACAA
6BF ACTTTAACTGCAAACGATGGTAGTTCGTTGGTGAATAATGTCCCTACTGGGGGACCACAA
6BSP ACTTTAACTGCAAACGATGGTAGTTCGTTGGTGAATAATGTCCCTACTGGGGGACCACAA
19AH ACTTTAACTGCAAACGATGGTAGTTCGTTGGTGAATAATGTCCCTACTGGGGGACCACAA
23FPO ACTTTAACTGCAAACGATGGTAGTTCGTTGGTGAATAATGTCCCTACTGGGGGACCACAA
19FTW ACTTTAACTGCAAACGATGGTAGTTCGTTGGTGAATAATGTCCCTACTGGGGGACCACAA
9VSP ACTTTAACTGCAAACGATGGTAGTTCGTTGGTGAATAATGTCCCTACTGGGGGACCACAA
TIGR4 ACTTTAACTGCAAACGATGGTAGTTCGTTGGTGAATAATGTCCCTACTGGGGGACCACAA
23FTW ACTTTAACTGCAAACGATGGTAGTTCGTTGGTGAATAATGTCCCTACTGGGGGACCACAA

Figure 196N

14CSR AATGATGGTGGCTTGCTAAAAAATGCAAAAGTGTCTATGATACGACTGAGAAAAGGATT
670 AATGATGGTGGCTTGCTAAAAAATGCAAAAGTGTCTATGATACGACTGAGAAAAGGATT
6BF AATGATGGTGGCTTGCTAAAAAATGCAAAAGTGTCTATGATACGACTGAGAAAAGGATT
6BSP AATGATGGTGGCTTGCTAAAAAATGCAAAAGTGTCTATGATACGACTGAGAAAAGGATT
19AH AATGATGGTGGCTTGCTAAAAAATGCAAAAGTGTCTATGATACGACTGAGAAAAGGATT
23FPO AATGATGGTGGCTTGCTAAAAAATGCAAAAGTGTCTATGATACGACTGAGAAAAGGATT
19FTW AATGATGGTGGCTTGCTAAAAAATGCAAAAGTGTCTATGATACGACTGAGAAAAGGATT
9VSP AATGATGGTGGCTTGCTAAAAAATGCAAAAGTGTCTATGATACGACTGAGAAAAGGATT
TIGR4 AATGATGGTGGTTTGTAAAAAATGCAAAAGTGTCTATGATACGACTGAGAAAAGGATT
23FTW AATGATGGTGGTTTGTAAAAAATGCAAAAGTGTCTATGATACGACTGAGAAAAGGATT

14CSR CGTGTAACAGGTTTGTACCTTGAACGGGTGAAAAAGTTACATTGACTTATAATGTTTCGC
670 CGTGTAACAGGTTTGTACCTTGAACGGGTGAAAAAGTTACATTGACTTATAATGTTTCGC
6BF CGTGTAACAGGTTTGTACCTTGAACGGGTGAAAAAGTTACATTGACTTATAATGTTTCGC
6BSP CGTGTAACAGGTTTGTACCTTGAACGGGTGAAAAAGTTACATTGACTTATAATGTTTCGC
19AH CGTGTAACAGGTTTGTACCTTGAACGGGTGAAAAAGTTACATTGACTTATAATGTTTCGC
23FPO CGTGTAACAGGTTTGTACCTTGAACGGGTGAAAAAGTTACATTGACTTATAATGTTTCGC
19FTW CGTGTAACAGGTTTGTACCTTGAACGGGTGAAAAAGTTACATTGACTTATAATGTTTCGC
9VSP CGTGTAACAGGTTTGTACCTTGAACGGGTGAAAAAGTTACATTGACTTATAATGTTTCGC
TIGR4 CGTGTAACAGGTTCTGTACCTTGAACGGGTGAAAAAGTTACGTTGACCTACAATGTTTCGT
23FTW CGTGTAACAGGTTCTGTACCTTGAACGGGTGAAAAAGTTACGTTGACCTACAATGTTTCGT

14CSR TTGAATGACCAATTTGTAAGCAATAAATTCTATGACACGAATGGTCGAACAACCCCTACAC
670 TTGAATGACCAATTTGTAAGCAATAAATTCTATGACACGAATGGTCGAACAACCCCTACAC
6BF TTGAATGACCAATTTGTAAGCAATAAATTCTATGACACGAATGGTCGAACAACCCCTACAC
6BSP TTGAATGACCAATTTGTAAGCAATAAATTCTATGACACGAATGGTCGAACAACCCCTACAC
19AH TTGAATGACCAATTTGTAAGCAATAAATTCTATGACACGAATGGTCGAACAACCCCTACAC
23FPO TTGAATGACCAATTTGTAAGCAATAAATTCTATGACACGAATGGTCGAACAACCCCTACAC
19FTW TTGAATGACCAATTTGTAAGCAATAAATTCTATGACACGAATGGTCGAACAACCCCTACAC
9VSP TTGAATGACCAATTTGTAAGCAATAAATTCTATGACACGAATGGTCGAACAACCCCTACAC
TIGR4 TTGAATGATGAGTTTGTAGCAATAAATTTTATGATACCAATGGTCGAACAACCCCTACAT
23FTW TTGAATGATGAGTTTGTAGCAATAAATTTTATGATACCAATGGTCGAACAACCCCTACAT

14CSR CCTAAGGAAGTAGAAAAGAACACAGTGC GCGACTTCCCGATTCCCTAAGATTTCGTGATGTA
670 CCTAAGGAAGTAGAAAAGAACACAGTGC GCGACTTCCCGATTCCCTAAGATTTCGTGATGTA
6BF CCTAAGGAAGTAGAAAAGAACACAGTGC GCGACTTCCCGATTCCCTAAGATTTCGTGATGTA
6BSP CCTAAGGAAGTAGAAAAGAACACAGTGC GCGACTTCCCGATTCCCTAAGATTTCGTGATGTA
19AH CCTAAGGAAGTAGAAAAGAACACAGTGC GCGACTTCCCGATTCCCTAAGATTTCGTGATGTA
23FPO CCTAAGGAAGTAGAAAAGAACACAGTGC GCGACTTCCCGATTCCCTAAGATTTCGTGATGTA
19FTW CCTAAGGAAGTAGAAAAGAACACAGTGC GCGACTTCCCGATTCCCTAAGATTTCGTGATGTA
9VSP CCTAAGGAAGTAGAAAAGAACACAGTGC GCGACTTCCCGATTCCCTAAGATTTCGTGATGTA
TIGR4 CCTAAGGAAGTAGAACAGAACACAGTGC GCGACTTCCCGATTCCCTAAGATTTCGTGATGTA
23FTW CCTAAGGAAGTAGAACAGAACACAGTGC GCGACTTCCCGATTCCCTAAGATTTCGTGATGTA

14CSR CGAAAGTATCCAGAAATCACAATTC AAAAGAGAAAAAACTTGGTGAAATTGAGTTTATT
670 CGAAAGTATCCAGAAATCACAATTC AAAAGAGAAAAAACTTGGTGAAATTGAGTTTATT
6BF CGAAAGTATCCAGAAATCACAATTC AAAAGAGAAAAAACTTGGTGAAATTGAGTTTATT
6BSP CGAAAGTATCCAGAAATCACAATTC AAAAGAGAAAAAACTTGGTGAAATTGAGTTTATT
19AH CGAAAGTATCCAGAAATCACAATTC AAAAGAGAAAAAACTTGGTGAAATTGAGTTTATT
23FPO CGAAAGTATCCAGAAATCACAATTC AAAAGAGAAAAAACTTGGTGAAATTGAGTTTATT
19FTW CGAAATATCCAGCAATTACGATTGCAAAAGAGAAAAAACTTGGTGAAATTGAGTTTATT
9VSP CGAAATATCCAGCAATTACGATTGCAAAAGAGAAAAAACTTGGTGAAATTGAGTTTATT
TIGR4 CGGAAGTATCCAGAAATCACAATTC AAAAGAGAAAAAACTTGGTGACATTGAGTTTATT
23FTW CGGAAGTATCCAGAAATCACAATTC AAAAGAGAAAAAACTTGGTGACATTGAGTTTATT

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14CSR AAGATCAATAAGAATGATAAAAAACCACTGAGAGATGCGGTCTTTAGTCTTCAAAAACAA
670 AAGATCAATAAGAATGATAAAAAACCACTGAGAGATGCGGTCTTTAGTCTTCAAAAACAA
6BF AAGATCAATAAGAATGATAAAAAACCACTGAGAGATGCGGTCTTTAGTCTTCAAAAACAA
6BSP AAGATCAATAAGAATGATAAAAAACCACTGAGAGATGCGGTCTTTAGTCTTCAAAAACAA
19AH AAGATCAATAAGAATGATAAAAAACCACTGAGAGATGCGGTCTTTAGTCTTCAAAAACAA
23FPO AAGATCAATAAGAATGATAAAAAACCACTGAGAGATGCGGTCTTTAGTCTTCAAAAACAA
19FTW AAGATCAATAAGAATGATAAAAAACCACTGAGAGATGCGGTCTTTAGTCTTCAAAAACAA
9VSP AAGATCAATAAGAATGATAAAAAACCACTGAGAGATGCGGTCTTTAGTCTTCAAAAACAA
TIGR4 AAGGTCAATAAAAAATGATAAAAAACCACTGAGAGGTGCGGTCTTTAGTCTTCAAAAACAA
23FTW AAGGTCAATAAAAAATGATAAAAAACCACTGAGAGATGCGGTCTTTAGTCTTCAAAAACAA
*** **

14CSR CATCCGGATTATCCAGATATTTATGGAGCTATTGATCAAAATGGCACTTATCAAAATGTG
670 CATCCGGATTATCCAGATATTTATGGAGCTATTGATCAAAATGGCACTTATCAAAATGTG
6BF CATCCGGATTATCCAGATATTTATGGAGCTATTGATCAAAATGGCACTTATCAAAATGTG
6BSP CATCCGGATTATCCAGATATTTATGGAGCTATTGATCAAAATGGCACTTATCAAAATGTG
19AH CATCCGGATTATCCAGATATTTATGGAGCTATTGATCAAAATGGCACTTATCAAAATGTG
23FPO CATCCGGATTATCCAGATATTTATGGAGCTATTGATCAAAATGGCACTTATCAAAATGTG
19FTW CATCCGGATTATCCAGATATTTATGGAGCTATTGATCAAAATGGCACTTATCAAAATGTG
9VSP CATCCGGATTATCCAGATATTTATGGAGCTATTGATCAAAATGGCACTTATCAAAATGTG
TIGR4 CATCCGGATTATCCAGATATTTATGGAGCTATTGATCAAAATGGCACTTATCAAAATGTG
23FTW CATCCGGATTATCCAGATATTTATGGAGCTATTGATCAAAATGGCACTTATCAAAATGTG

14CSR AGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGATGGGAAATATCGATTA
670 AGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGATGGGAAATATCGATTA
6BF AGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGATGGGAAATATCGATTA
6BSP AGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGATGGGAAATATCGATTA
19AH AGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGATGGGAAATATCGATTA
23FPO AGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGATGGGAAATATCGATTA
19FTW AGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGATGGGAAATATCGATTA
9VSP AGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGATGGGAAATATCGATTA
TIGR4 AGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGATGGGAAATATCGATTA
23FTW AGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTGATGGGAAATATCGATTA

14CSR TTTGAAAATTCTGAACAGCTGGTTATAAACCCGTTCAAATAAGCCTATCGTTGCCTTC
670 TTTGAAAATTCTGAACAGCTGGTTATAAACCCGTTCAAATAAGCCTATCGTTGCCTTC
6BF TTTGAAAATTCTGAACAGCTGGTTATAAACCCGTTCAAATAAGCCTATCGTTGCCTTC
6BSP TTTGAAAATTCTGAACAGCTGGTTATAAACCCGTTCAAATAAGCCTATCGTTGCCTTC
19AH TTTGAAAATTCTGAACAGCTGGTTATAAACCCGTTCAAATAAGCCTATCGTTGCCTTC
23FPO TTTGAAAATTCTGAACAGCTGGTTATAAACCCGTTCAAATAAGCCTATCGTTGCCTTC
19FTW TTTGAAAATTCTGAACAGCTGGTTATAAACCCGTTCAAATAAGCCTATCGTTGCCTTC
9VSP TTTGAAAATTCTGAACAGCTGGTTATAAACCCGTTCAAATAAGCCTATCGTTGCCTTC
TIGR4 TTTGAAAATTCTGAACAGCTGGTTATAAACCCGTTCAAATAAGCCTATCGTTGCCTTC
23FTW TTTGAAAATTCTGAACAGCTGGTTATAAACCCGTTCAAATAAGCCTATCGTTGCCTTC

14CSR CAAATAGTAAATGGAGAAGTCAGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCG
670 CAAATAGTAAATGGAGAAGTCAGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCG
6BF CAAATAGTAAATGGAGAAGTCAGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCG
6BSP CAAATAGTAAATGGAGAAGTCAGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCG
19AH CAAATAGTAAATGGAGAAGTCAGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCG
23FPO CAAATAGTAAATGGAGAAGTCAGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCG
19FTW CAAATAGTAAATGGAGAAGTCAGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCG
9VSP CAAATAGTAAATGGAGAAGTCAGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCG
TIGR4 CAAATAGTAAATGGAGAAGTCAGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCG
23FTW CAAATAGTAAATGGAGAAGTCAGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCG

Figure 196P

PCT/US05/27239/409/487

14CSR GGTTACGAGTTTACGAATGATAAGCACTATATCACAAATGAGCCAATTCTCTCCAAAAAGA
670 GGTTACGAGTTTACGAATGATAAGCACTATATCACAAATGAGCCAATTCTCTCCAAAAAGA
6BF GGTTACGAGTTTACGAATGATAAGCACTATATCACAAATGAGCCAATTCTCTCCAAAAAGA
6BSP GGTTACGAGTTTACGAATGATAAGCACTATATCACAAATGAGCCAATTCTCTCCAAAAAGA
19AH GGTTACGAGTTTACGAATGATAAGCACTATATCACAAATGAGCCAATTCTCTCCAAAAAGA
23FPO GGTTACGAGTTTACGAATGATAAGCACTATATCACAAATGAGCCAATTCTCTCCAAAAAGA
19FTW GGTTACGAGTTTACGAATGATAAGCACTATATTACCAATGAACCTATTCTCTCCAAAGAGA
9VSP GGTTACGAGTTTACGAATGATAAGCACTATATTACCAATGAACCTATTCTCTCCAAAGAGA
TIGR4 GGTTACGAGTTTACGAATGATAAGCACTATATTACCAATGAACCTATTCTCTCCAAAGAGA
23FTW GGTTACGAGTTTACGAATGATAAGCACTATATTACCAATGAACCTATTCTCTCCAAAGAGA
***** ** ***** ** ***** **

14CSR GAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATG
670 GAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATG
6BF GAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATG
6BSP GAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATG
19AH GAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATG
23FPO GAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATG
19FTW GAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATG
9VSP GAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATG
TIGR4 GAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATG
23FTW GAATATCCTCGAAGTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATG
***** *****

14CSR ATGGGAGGAGTTCTATTATACACACGGAACATCCGTAAAGTGTAGCAATGAGAAATGAT
670 ATGGGAGGAGTTCTATTATACACACGGAACATCCGTAAAGTGTAGCAATGAGAAATGAT
6BF ATGGGAGGAGTTCTATTATACACACGGAACATCCGTAAAGTGTAGCAATGAGAAATGAT
6BSP ATGGGAGGAGTTCTATTATACACACGGAACATCCGTAAAGTGTAGCAATGAGAAATGAT
19AH ATGGGAGGAGTTCTATTATACACACGGAACATCCGTAAAGTGTAGCAATGAGAAATGAT
23FPO ATGGGAGGAGTTCTATTATACACACGGAACATCCGTAAAGTGTAGCAATGAGAAATGAT
19FTW ATGGGAGGAGTTCTATTATACACACGGAACATCCGTAAAGTGTAG-----AAATGAT
9VSP ATGGGAGGAGTTCTATTATACACACGGAACATCCGTAAAGTGTAG-----AAATGAT
TIGR4 ATGGGAGGAGTTCTATTATACACACGGAACATCCGTAAAGTGTAG-----AAATGAT
23FTW ATGGGAGGAGTTCTATTATACACACGGAACATCCGTAAAGTGTAG-----AAATGAT
***** *****

14CSR AATATCGATACTCTGAGCGATACTTTTAAAGAAGTAGCACTCAAGAAGAGATTTAAGTTTA
670 AATATCGATACTCTGAGCGATACTTTTAAAGAAGTAGCACTCAAGAAGAGATTTAAGTTTA
6BF AATATCGATACTCTGAGCGATACTTTTAAAGAAGTAGCACTCAAGAAGAGATTTAAGTTTA
6BSP AATATCGATACTCTGAGCGATACTTTTAAAGAAGTAGCACTCAAGAAGAGATTTAAGTTTA
19AH AATATCGATACTCTGAGCGATACTTTTAAAGAAGTAGCACTCAAGAAGAGATTTAAGTTTA
23FPO AATATCGATACTCTGAGCGATACTTTTAAAGAAGTAGCACTCAAGAAGAGATTTAAGTTTA
19FTW AATATCTATGTTCTGAACAATACTTTTAAAGAAGTAGCACTCAAGAAGAGATTTAAGTTTA
9VSP AATATCTATGTTCTGAACGATACTTTTAAAGAAGTAGCACTCAAGAAGAGATTTAAGTTTA
TIGR4 AATATCTATGTTCTGAACGATACTTTTAAAGAAGTAGCACTCAAGAAGAGATTTAAGTTTA
23FTW AATATCTATGTTCTGAACGATACTTTTAAAGAAGTAGCACTCAAGAAGAGATTTAAGTTTA
***** ** ***** * *****

14CSR CTTGGTGAAAACAGTTTCTTCGCCAAGTAAACCACCATTGAAAGGGGAGATGTTTTCGA
670 CTTGGTGAAAACAGTTTCTTCGCCAAGTAAACCACCATTGAAAGGGGAGATGTTTTCGA
6BF CTTGGTGAAAACAGTTTCTTCGCCAAGTAAACCACCATTGAAAGGGGAGATGTTTTCGA
6BSP CTTGGTGAAAACAGTTTCTTCGCCAAGTAAACCACCATTGAAAGGGGAGATGTTTTCGA
19AH CTTGGTGAAAACAGTTTCTTCGCCAAGTAAACCACCATTGAAAGGGGAGATGTTTTCGA
23FPO CTTGGTGAAAACAGTTTCTTCGCCAAGTAAACCACCATTGAAAGGGGAGATGTTTTCGA
19FTW CTTGGTGAAAACAGTTTCTTCGCCAAGTAAACCACCATTGAAAGGGGAGATGTTTTCGA
9VSP CTTGGTGAAAACAGTTTCTTCGCCAAGTAAACCACCATTGAAAGGGGAGATGTTTTCGA
TIGR4 CTTGGTGAAAACAGTTTCTTCGCCAAGTAAACCACCATTGAAAGGGGAGATGTTTTCGA
23FTW CTTGGTGAAAACAGTTTCTTCGCCAAGTAAACCACCATTGAAAGGGGAGATGTTTTCGA
***** * ***** * *****

14CSR AAACCTTGACAGAAAAAAGGATTATTATTGTCATGTGTAATTCATTACATTGCTCACAGT
670 AAACCTTGACAGAAAAAAGGATTATTATTGTCATGTGTAATTCATTACATTGCTCACAGT
6BF AAACCTTGACAGAAAAAAGGATTATTATTGTCATGTGTAATTCATTACATTGCTCACAGT
6BSP AAACCTTGACAGAAAAAAGGATTATTATTGTCATGTGTAATTCATTACATTGCTCACAGT
19AH AAACCTTGACAGAAAAAAGGATTATTATTGTCATGTGTAATTCATTACATTGCTCACAGT
23FPO AAACCTTGACAGAAAAAAGGATTATTATTGTCATGTGTAATTCATTACATTGCTCACAGT
19FTW AAACCTTGACAGAAAAA--GGATTATTATTGTCATGTGTAATTCATTACATTGCTCACAGT
9VSP AAACCTTGACAGAAAAA--GGATTATTATTGTCATGTGTAATTCATTACATTGCTCACAGT
TIGR4 AAACCTTGACAGAAAAA--GGATTATTATTGTCATGTGTAATTCATTACATTGCTCACAGT
23FTW AAACCTTGACAGAAAAA--GGATTATTATTGTCATGTGTAATTCATTACATTGCTCACAGT

14CSR TGATTTTAAGAGATATGAATAAGGAGAAATCATGAAATCAATCAACAAATTTTAAACAAT
670 TGATTTTAAGAGATATGAATAAGGAGAAATCATGAAATCAATCAACAAATTTTAAACAAT
6BF TGATTTTAAGAGATATGAATAAGGAGAAATCATGAAATCAATCAACAAATTTTAAACAAT
6BSP TGATTTTAAGAGATATGAATAAGGAGAAATCATGAAATCAATCAACAAATTTTAAACAAT
19AH TGATTTTAAGAGATATGAATAAGGAGAAATCATGAAATCAATCAACAAATTTTAAACAAT
23FPO TGATTTTAAGAGATATGAATAAGGAGAAATCATGAAATCAATCAACAAATTTTAAACAAT
19FTW TGATTTTAAGAGATA--AATAAGGAGAAATCATGAAATCAATCAACAAATTTTAAACAAT
9VSP TGATTTTAAGAGATA--AATAAGGAGAAATCATGAAATCAATCAACAAATTTTAAACAAT
TIGR4 TGATTTTAAGAGATATGAATAAGGAGAAATCATGAAATCAATCAACAAATTTTAAACAAT
23FTW TGATTTTAAGAGATATGAATAAGGAGAAATCATGAAATCAATCAACAAATTTTAAACAAT

14CSR GCTTGCTGCCTTATTACTGACAGCGAGTAGCCTGTTTTGCTGCAACAGTTTTTGC GGC
670 GCTTGCTGCCTTATTACTGACAGCGAGTAGCCTGTTTTGCTGCAACAGTTTTTGC GGC
6BF GCTTGCTGCCTTATTACTGACAGCGAGTAGCCTGTTTTGCTGCAACAGTTTTTGC GGC
6BSP GCTTGCTGCCTTATTACTGACAGCGAGTAGCCTGTTTTGCTGCAACAGTTTTTGC GGC
19AH GCTTGCTGCCTTATTACTGACAGCGAGTAGCCTGTTTTGCTGCAACAGTTTTTGC GGC
23FPO GCTTGCTGCCTTATTACTGACAGCGAGTAGCCTGTTTTGCTGCAACAGTTTTTGC GGC
19FTW GCTTGCTGCCTTATTATTGACAGCGAGTAGCCTGTTTTGCTGCAACAGTTTTTGC GGC
9VSP GCTTGCTGCCTTATTACTGACAGCGAGTAGCCTGTTTTGCTGCAACAGTTTTTGC GGC
TIGR4 GCTTGCTGCCTTATTACTGACAGCGAGTAGCCTGTTTTGCTGCAACAGTTTTTGC GGC
23FTW ACTTGCTGCCTTATTACTGACAGCGAGTAGCCTGTTCTGCTGCAACAGTTTTTGC GGC

14CSR GGACAATGTTAGTACAGCACCAGATGCTGTTACTAAAACCTTAAACAATCCATAAGTTACT
670 GGACAATGTTAGTACAGCACCAGATGCTGTTACTAAAACCTTAAACAATCCATAAGTTACT
6BF GGACAATGTTAGTACAGCACCAGATGCTGTTACTAAAACCTTAAACAATCCATAAGTTACT
6BSP GGACAATGTTAGTACAGCACCAGATGCTGTTACTAAAACCTTAAACAATCCATAAGTTACT
19AH GGACAATGTTAGTACAGCACCAGATGCTGTTACTAAAACCTTAAACAATCCATAAGTTACT
23FPO GGACAATGTTAGTACAGCACCAGATGCTGTTACTAAAACCTTAAACAATCCATAAGTTACT
19FTW TGG-GACGACA--ACAACATCTGTTACCGTTCATAAACTATTGGCAACAGATGGGGATAT
9VSP TGG-GACGACA--ACAACATCTGTTACCGTTCATAAACTATTGGCAACAGATGGGGATAT
TIGR4 TGG-GACGACA--ACAACATCTGTTACCGTTCATAAACTATTGGCAACAGATGGGGATAT
23FTW GGA-ACAAAA--ACTAAGCACTTACAGTTACATAAATATTGATGACAGATCAAGAGCT
* * * * *

14CSR GCTCTCA---GAAGATGATTTAAAGACTTGGGATACAAACGGTCCTAA--AGGATATGATG
670 GCTCTCA---GAAGATGATTTAAAGACTTGGGATACAAACGGTCCTAA--AGGATATGATG
6BF GCTCTCA---GAAGATGATTTAAAGACTTGGGATACAAACGGTCCTAA--AGGATATGATG
6BSP GCTCTCA---GAAGATGATTTAAAGACTTGGGATACAAACGGTCCTAA--AGGATATGATG
19AH GCTCTCA---GAAGATGATTTAAAGACTTGGGATACAAACGGTCCTAA--AGGATATGATG
23FPO GCTCTCA---GAAGATGATTTAAAGACTTGGGATACAAACGGTCCTAA--AGGATATGATG
19FTW GGATAAAATTGCAATGAGTTAGAAACAGGTAACCTATGCTGGTAATAA--AGTGGGTGTTTC
9VSP GGATAAAATTGCAATGAGTTAGAAACAGGTAACCTATGCTGGTAATAA--AGTGGGTGTTTC
TIGR4 GGATAAAATTGCAATGAGTTAGAAACAGGTAACCTATGCTGGTAATAA--AGTGGGTGTTTC
23FTW TGAC-----GCTTGAATTCTGATGCGATTACTACTGCAGGTTATGACGGTTCGCAAAA
* * * * *

Figure 196R

14CSR GAACTCAATCTAGTTTAAAAGATTTAACTGGAGTTGTAGCTG----AGGAAATTCCAAAT
670 GAACTCAATCTAGTTTAAAAGATTTAACTGGAGTTGTAGCTG----AGGAAATTCCAAAT
6BF GAACTCAATCTAGTTTAAAAGATTTAACTGGAGTTGTAGCTG----AGGAAATTCCAAAT
6BSP GAACTCAATCTAGTTTAAAAGATTTAACTGGAGTTGTAGCTG----AGGAAATTCCAAAT
19AH GAACTCAATCTAGTTTAAAAGATTTAACTGGAGTTGTAGCTG----AGGAAATTCCAAAT
23FPO GAACTCAATCTAGTTTAAAAGATTTAACTGGAGTTGTAGCTG----AGGAAATTCCAAAT
19FTW TACCTGCA---AATGCAAAAGAAATTGCCGGTGTATGTTTCGTTTGGACAAATACTAATA
9VSP TACCTGCA---AATGCAAAAGAAATTGCCGGTGTATGTTTCGTTTGGACAAATACTAATA
TIGR4 TACCTGCA---AATGCAAAAGAAATTGCCGGTGTATGTTTCGTTTGGACAAATACTAATA
23FTW T-TTGAA---CAGTTCAAACAACCTCAAGGTGTTCCACAAG---GAGTAACCGAAATCT
* * * * * * * * * *

14CSR GTATACTT-----TGAATTACAAAAGTATA-ATTTGACTGATGGT--AAGGAAAAAGA
670 GTATACTT-----TGAATTACAAAAGTATA-ATTTGACTGATGGT--AAGGAAAAAGA
6BF GTATACTT-----TGAATTACAAAAGTATA-ATTTGACTGATGGT--AAGGAAAAAGA
6BSP GTATACTT-----TGAATTACAAAAGTATA-ATTTGACTGATGGT--AAGGAAAAAGA
19AH GTATACTT-----TGAATTACAAAAGTATA-ATTTGACTGATGGT--AAGGAAAAAGA
23FPO GTATACTT-----TGAATTACAAAAGTATA-ATTTGACTGATGGT--AAGGAAAAAGA
19FTW ATGAAATTATTGATGAAAATGGCCAACTCTAGGAGTGAATATTGATCCACAAACATTTA
9VSP ATGAAATTATTGATGAAAATGGCCAACTCTAGGAGTGAATATTGATCCACAAACATTTA
TIGR4 ATGAAATTATTGATGAAAATGGCCAACTCTAGGAGTGAATATTGATCCACAAACATTTA
23FTW CTGGTGTTCG--ATTTCGAGTTACAGAGTTATACGGGTCTCAAGGA--AAAGAACAAGAA
* ** * * * * * * * *

14CSR AAATCTTAAAGATGATAGTAAATGGACAACAGTTCATGGTGGTTTGACAACATAAGATGG
670 AAATCTTAAAGATGATAGTAAATGGACAACAGTTCATGGTGGTTTGACAACATAAGATGG
6BF AAATCTTAAAGATGATAGTAAATGGACAACAGTTCATGGTGGTTTGACAACATAAGATGG
6BSP AAATCTTAAAGATGATAGTAAATGGACAACAGTTCATGGTGGTTTGACAACATAAGATGG
19AH AAATCTTAAAGATGATAGTAAATGGACAACAGTTCATGGTGGTTTGACAACATAAGATGG
23FPO AAATCTTAAAGATGATAGTAAATGGACAACAGTTCATGGTGGTTTGACAACATAAGATGG
19FTW AACTCTCAGGGGCAATGCCGGC--AACTGCAATGAAAAAATTAACAGAAGCTGAA---GG
9VSP AACTCTCAGGGGCAATGCCGGC--AACTGCAATGAAAAAATTAACAGAAGCTGAA---GG
TIGR4 AACTCTCAGGGGCAATGCCGGC--AACTGCAATGAAAAAATTAACAGAAGCTGAA---GG
23FTW AA-TTTAACGAATGATGCGGTTTGGACTGCGGTTAATAAAGGTGTGACAGTGAACAGG
* * * * * * * * *

14CSR ACTTAAAAATTGAAACCAGTACTCTTAAAGGTGT---GTATCGTATTCGTGAGGATAGAAC
670 ACTTAAAAATTGAAACCAGTACTCTTAAAGGTGT---GTATCGTATTCGTGAGGATAGAAC
6BF ACTTAAAAATTGAAACCAGTACTCTTAAAGGTGT---GTATCGTATTCGTGAGGATAGAAC
6BSP ACTTAAAAATTGAAACCAGTACTCTTAAAGGTGT---GTATCGTATTCGTGAGGATAGAAC
19AH ACTTAAAAATTGAAACCAGTACTCTTAAAGGTGT---GTATCGTATTCGTGAGGATAGAAC
23FPO ACTTAAAAATTGAAACCAGTACTCTTAAAGGTGT---GTATCGTATTCGTGAGGATAGAAC
19FTW AGCTAAATTTAACACGGCAAATTTACCAGCTGCTAAGTATAAAATTTATGAAATTCACAG
9VSP AGCTAAATTTAACACGGCAAATTTACCAGCTGCTAAGTATAAAATTTATGAAATTCACAG
TIGR4 AGCTAAATTTAACACGGCAAATTTACCAGCTGCTAAGTATAAAATTTATGAAATTCACAG
23FTW TGTTAAATTTGATACTGAAGTTTACAAGGGAC---ATATCGTCTGTGCAAGTACGTAA
* * * * * * * * *

14CSR AAAGACTACCTATGTTGGTCCCTAATGGGCAAGTATTAACAGGTTCAAAGCCGTACCTGC
670 AAAGACTACCTATGTTGGTCCCTAATGGGCAAGTATTAACAGGTTCAAAGCCGTACCTGC
6BF AAAGACTACCTATGTTGGTCCCTAATGGGCAAGTATTAACAGGTTCAAAGCCGTACCTGC
6BSP AAAGACTACCTATGTTGGTCCCTAATGGGCAAGTATTAACAGGTTCAAAGCCGTACCTGC
19AH AAAGACTACCTATGTTGGTCCCTAATGGGCAAGTATTAACAGGTTCAAAGCCGTACCTGC
23FPO AAAGACTACCTATGTTGGTCCCTAATGGGCAAGTATTAACAGGTTCAAAGCCGTACCTGC
19FTW TTTATCAACTTATGTCGGTGAAGATGGAGCAACCTTAACAGGTTCTAAAGCAGTTCCAAT
9VSP TTTATCAACTTATGTCGGTGAAGATGGAGCAACCTTAACAGGTTCTAAAGCAGTTCCAAT
TIGR4 TTTATCAACTTATGTCGGTGAAGATGGAGCAACCTTAACAGGTTCTAAAGCAGTTCCAAT
23FTW AGAATCGACTTATGTCGGTCCAAATGGTAAAGTTTAAACAGGTATGAAAGCTGTTCTCTGC
* * * * * * * * *

Figure 196S

PCT/US05/27239 412/487

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14CSR      TCTTGTAACCTCTTCCACTTGTTAACAATAATGGTACAGTAATTGATGCACATGTTTTCCC
670        TCTTGTAACCTCTTCCACTTGTTAACAATAATGGTACAGTAATTGATGCACATGTTTTCCC
6BF        TCTTGTAACCTCTTCCACTTGTTAACAATAATGGTACAGTAATTGATGCACATGTTTTCCC
6BSP       TCTTGTAACCTCTTCCACTTGTTAACAATAATGGTACAGTAATTGATGCACATGTTTTCCC
19AH       TCTTGTAACCTCTTCCACTTGTTAACAATAATGGTACAGTAATTGATGCACATGTTTTCCC
23FPO      TCTTGTAACCTCTTCCACTTGTTAACAATAATGGTACAGTAATTGATGCACATGTTTTCCC
19FTW      TGAAATTGAATTACCATT-----GAACGATGTTGTGGA---TGCGCATGTGTATCC
9VSP       TGAAATTGAATTACCATT-----GAACGATGTTGTGGA---TGCGCATGTGTATCC
TIGR4      TGAAATTGAATTACCATT-----GAACGATGTTGTGGA---TGCGCATGTGTATCC
23FTW      TTTAATTACTCTGCCGCTTGTAACCAAAATGGTGTGTAGAAAATGCACATGCTATCC
          *      *      *      *      *      *      *      *      *      *
14CSR      TAAAAATTCATATAATAAACCAGTTGTAGATAAAAGAATTGCTGATACTTTGAATTATAA
670        TAAAAATTCATATAATAAACCAGTTGTAGATAAAAGAATTGCTGATACTTTGAATTATAA
6BF        TAAAAATTCATATAATAAACCAGTTGTAGATAAAAGAATTGCTGATACTTTGAATTATAA
6BSP       TAAAAATTCATATAATAAACCAGTTGTAGATAAAAGAATTGCTGATACTTTGAATTATAA
19AH       TAAAAATTCATATAATAAACCAGTTGTAGATAAAAGAATTGCTGATACTTTGAATTATAA
23FPO      TAAAAATTCATATAATAAACCAGTTGTAGATAAAAGAATTGCTGATACTTTGAATTATAA
19FTW      AAAAAATACAGAAGCAAAGCCAAAAATTGATAAAGATTTCAAAGGTAAAGCAAATCCAGA
9VSP       AAAAAATACAGAAGCAAAGCCAAAAATTGATAAAGATTTCAAAGGTAAAGCAAATCCAGA
TIGR4      AAAAAATACAGAAGCAAAGCCAAAAATTGATAAAGATTTCAAAGGTAAAGCAAATCCAGA
23FTW      AAAGAATTCTGAAGACAACCTACAGCAACGAAACATTTGATACTGCAGCAGGTTTCGT
          **  ***  *      *      *      *      *      *      *      *
14CSR      CGATCAA-----AATGGTCTGTCTATCGGTACTAAAATCCCATATGTTGT----TA
670        CGATCAA-----AATGGTCTGTCTATCGGTACTAAAATCCCATATGTTGT----TA
6BF        CGATCAA-----AATGGTCTGTCTATCGGTACTAAAATCCCATATGTTGT----TA
6BSP       CGATCAA-----AATGGTCTGTCTATCGGTACTAAAATCCCATATGTTGT----TA
19AH       CGATCAA-----AATGGTCTGTCTATCGGTACTAAAATCCCATATGTTGT----TA
23FPO      CGATCAA-----AATGGTCTGTCTATCGGTACTAAAATCCCATATGTTGT----TA
19FTW      TACACCACGTGTAGATAAAGATACACCTGTGAACCACCAAGTTGGAGATGTTGTAGAGTA
9VSP       TACACCACGTGTAGATAAAGATACACCTGTGAACCACCAAGTTGGAGATGTTGTAGAGTA
TIGR4      TACACCACGTGTAGATAAAGATACACCTGTGAACCACCAAGTTGGAGATGTTGTAGAGTA
23FTW      AGATCCAGGTG---AAAAAGGTTTAGCAATTGGCACTAAGGTACCGTATATTGT----TA
          *      *      *      *      *      *      *      *      *      *
14CSR      ATACAACAATTCCAAGTAATGCAACATT-----TGCAACTTCATTTTGGTCAGATG
670        ATACAACAATTCCAAGTAATGCAACATT-----TGCAACTTCATTTTGGTCAGATG
6BF        ATACAACAATTCCAAGTAATGCAACATT-----TGCAACTTCATTTTGGTCAGATG
6BSP       ATACAACAATTCCAAGTAATGCAACATT-----TGCAACTTCATTTTGGTCAGATG
19AH       ATACAACAATTCCAAGTAATGCAACATT-----TGCAACTTCATTTTGGTCAGATG
23FPO      ATACAACAATTCCAAGTAATGCAACATT-----TGCAACTTCATTTTGGTCAGATG
19FTW      CGA-AATTGTTACAAAAATCCAGCACTTGCTAATTATGCAACAGCAAACCTGGAGCGATA
9VSP       CGA-AATTGTTACAAAAATCCAGCACTTGCTAATTATGCAACAGCAAACCTGGAGCGATA
TIGR4      CGA-AATTGTTACAAAAATCCAGCACTTGCTAATTATGCAACAGCAAACCTGGAGCGATA
23FTW      CAACAACATATCCGAAAAACTCAACTCT-----TGCAACAGCTTCTGGTCAGATG
          *      *      *      *      *      *      *      *      *      *
14CSR      AAATGACAGAAGGTCTAACTTATAATGAAGA-GTAACAA---TTACTTTGAATAATGTAG
670        AAATGACAGAAGGTCTAACTTATAATGAAGATGTAACAA---TTACTTTGAATAATGTAG
6BF        AAATGACAGAAGGTCTAACTTATAATGAAGATGTAACAA---TTACTTTGAATAATGTAG
6BSP       AAATGACAGAAGGTCTAACTTATAATGAAGATGTAACAA---TTACTTTGAATAATGTAG
19AH       AAATGACAGAAGGTCTAACTTATAATGAAGATGTAACAA---TTACTTTGAATAATGTAG
23FPO      AAATGACAGAAGGTCTAACTTATAATGAAGATGTAACAA---TTACTTTGAATAATGTAG
19FTW      GAATGACTGAAGGTTTGGCATTCAACAAAGGTACAGTGAAAGTAACTGTTGATGATGTTG
9VSP       GAATGACTGAAGGTTTGGCATTCAACAAAGGTACAGTGAAAGTAACTGTTGATGATGTTG
TIGR4      GAATGACTGAAGGTTTGGCATTCAACAAAGGTACAGTGAAAGTAACTGTTGATGATGTTG
23FTW      AAATGACAGAAGGTCTAGATTATAATGGTGATGATT---GTTAATTATAATGGTCAAC
          *****  *****  *      *      *      *      *      *      *

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Figure 196T

414/487

14CSR CTCCAATCAATC-TGAAGAACCACGTTGTA AAAACATACGGTAAAAAGTTTGTCAAAGTAG
670 CTCCAATCAATCCTGAAGAACCACGTTGTA AAAACATACGGTAAAAAGTTTGTCAAAGTAG
6BF CTCCAATCAATCCTGAAGAACCACGTTGTA AAAACATACGGTAAAAAGTTTGTCAAAGTAG
6BSP CTCCAATCAATCCTGAAGAACCACGTTGTA AAAACATACGGTAAAAAGTTTGTCAAAGTAG
19AH CTCCAATCAATCCTGAAGAACCACGTTGTA AAAACATACGGTAAAAAGTTTGTCAAAGTAG
23FPO CTCCAATCAATCTTGAAGAACCACGTTGTA AAAACATACGGTAAAAAGTTTGTCAAAGTAG
19FTW AACCACCTTGATCCAACAGAGCCAAAAGTTGTTACATATGGTAAAAAGTTTGTCAAAGTTA
9VSP AACCACCTTGATCCAACAGAGCCAAAAGTTGTTACATATGGTAAAAAGTTTGTCAAAGTTA
TIGR4 AACCACCTTGATCCAACAGAGCCAAAAGTTGTTACATATGGTAAAAAGTTTGTCAAAGTTA
23FTW AACCACATAATCCTGAAGAGCCACGTTGTA AAAACATATGGTAAAAATTCGTTAAGGTTG
*** * *** **

Figure 196V

Figure 196W

PCT/US2005/027239 416/487

14CSR TCTTACTTCTAATACGGATGGTCAATTCCAAATTTTCAGGTCTTGCTGCTGGTACTTATAA
670 TCTTACTTCTAATACGGATGGTCAATTCCAAATTTTCAGGTCTTGCTGCTGGTACTTATAA
6BF TCTTACTTCTAATACGGATGGTCAATTCCAAATTTTCAGGTCTTGCTGCTGGTACTTATAA
6BSP TCTTACTTCTAATACGGATGGTCAATTCCAAATTTTCAGGTCTTGCTGCTGGTACTTATAA
19AH TCTTACTTCTAATACGGATGGTCAATTCCAAATTTTCAGGTCTTGCTGCTGGTACTTATAA
23FPO TCTTACTTCTAATACGGATGGTCAATTCCAAATTTTCAGGTCTTGCTGCTGGTACTTATAA
19FTW ATTAGTTTCTGATGCACAAGGTGCGCTTTGAAATTACAGGCCCTTCTGTCAGGTACATATTA
9VSP ATTAGTTTCTGATGCACAAGGTGCGCTTTGAAATTACAGGCCCTTCTGTCAGGTACATATTA
TIGR4 ATTAGTTTCTGATGCACAAGGTGCGCTTTGAAATTACAGGCCCTTCTGTCAGGTACATATTA
23FTW TCTTACTTCTAACACTGATGGTCAATTCCAAATTTTCAGGTCTTGCTGCTGGAAGCTACAC
* * * * *

14CSR ATTAGAAGAAATTAAAGCTCCAGAAGGTTTTGCGAAAAT---TGATGATGTAGAATTTGT
670 ATTAGAAGAAATTAAAGCTCCAGAAGGTTTTGCGAAAAT---TGATGATGTAGAATTTGT
6BF ATTAGAAGAAATTAAAGCTCCAGAAGGTTTTGCGAAAAT---TGATGATGTAGAATTTGT
6BSP ATTAGAAGAAATTAAAGCTCCAGAAGGTTTTGCGAAAAT---TGATGATGTAGAATTTGT
19AH ATTAGAAGAAATTAAAGCTCCAGAAGGTTTTGCGAAAAT---TGATGATGTAGAATTTGT
23FPO ATTAGAAGAAATTAAAGCTCCAGAAGGTTTTGCGAAAAT---TGATGATGTAGAATTTGT
19FTW CTTAGAAGAAACAAAACAGCCTGCTGGTTATGCATTACTAAGTCCGTCAGAAATTTGA
9VSP CTTAGAAGAAACAAAACAGCCTGCTGGTTATGCATTACTAAGTCCGTCAGAAATTTGA
TIGR4 CTTAGAAGAAACAAAACAGCCTGCTGGTTATGCATTACTAAGTCCGTCAGAAATTTGA
23FTW GTTGAAGAAACAAAAGCTCCAGAAGGTTTTGCAAACT---TGGAGATGTGAAGTTTGA
* * * * *

14CSR TGTTGGAGCAGGTTCTTG-----GAATCAAGGTGAGTTTAATTACTTAAAAGATGTTCA
670 TGTTGGAGCAGGTTCTTG-----GAATCAAGGTGAGTTTAATTACTTAAAAGATGTTCA
6BF TGTTGGAGCAGGTTCTTG-----GAATCAAGGTGAGTTTAATTACTTAAAAGATGTTCA
6BSP TGTTGGAGCAGGTTCTTG-----GAATCAAGGTGAGTTTAATTACTTAAAAGATGTTCA
19AH TGTTGGAGCAGGTTCTTG-----GAATCAAGGTGAGTTTAATTACTTAAAAGATGTTCA
23FPO TGTTGGAGCAGGTTCTTG-----GAATCAAGGTGAGTTTAATTACTTAAAAGATGTTCA
19FTW AGTCACTGCAACTTCTTATTCAGCGACTGGACAAGGCATTGAGTATACTGCTGGTTCAGG
9VSP AGTCACTGCAACTTCTTATTCAGCGACTGGACAAGGCATTGAGTATACTGCTGGTTCAGG
TIGR4 AGTCACTGCAACTTCTTATTCAGCGACTGGACAAGGCATTGAGTATACTGCTGGTTCAGG
23FTW GGTGGAGCAGGTTCTTG-----GAATCAAGGTGAGTTTAATTACTTAAAAGATGTTCA
* * * * *

14CSR AAAGAATGACGCTACAAAAGTAGTCAACAAAAAAATCACTATCCCACAAACGGGTGGTAT
670 AAAGAATGACGCTACAAAAGTAGTCAACAAAAAAATCACTATCCCACAAACGGGTGGTAT
6BF AAAGAATGACGCTACAAAAGTAGTCAACAAAAAAATCACTATCCCACAAACGGGTGGTAT
6BSP AAAGAATGACGCTACAAAAGTAGTCAACAAAAAAATCACTATCCCACAAACGGGTGGTAT
19AH AAAGAATGACGCTACAAAAGTAGTCAACAAAAAAATCACTATCCCACAAACGGGTGGTAT
23FPO AAAGAATGACGCTACAAAAGTAGTCAACAAAAAAATCAGATCCCACAAACGGGTGGTAT
19FTW TAAAGATGACGCTACAAAAGTAGTCAACAAAAAAATCAGATCCCACAAACGGGTGGTAT
9VSP TAAAGATGACGCTACAAAAGTAGTCAACAAAAAAATCAGATCCCACAAACGGGTGGTAT
TIGR4 TAAAGATGACGCTACAAAAGTAGTCAACAAAAAAATCACTATCCCACAAACGGGTGGTAT
23FTW GAAGAACGACGCTACAAAAGTAGTCAACAAAAAAATCAGATCCCTCAAACGGGTGGTAT
* * * * *

14CSR TGGTACAATTATCTTTGCTGTAGCGGGGGCTGCGATTATGGGTATTGCAGTGTACGCATA
670 TGGTACAATTATCTTTGCTGTAGCGGGGGCTGCGATTATGGGTATTGCAGTGTACGCATA
6BF TGGTACAATTATCTTTGCTGTAGCGGGGGCTGCGATTATGGGTATTGCAGTGTACGCATA
6BSP TGGTACAATTATCTTTGCTGTAGCGGGGGCTGCGATTATGGGTATTGCAGTGTACGCATA
19AH TGGTACAATTATCTTTGCTGTAGCGGGGGCTGCGATTATGGGTATTGCAGTGTACGCATA
23FPO TGGTACAATTATCTTTGCTGTAGCAGGGGGCTGTGATTATGGGTATTGCAGTGTACGCATA
19FTW TGGTACAATTATCTTTGCTGTAGCAGGGGGCTGTGATTATGGGTATTGCAGTGTACGCATA
9VSP TGGTACAATTATCTTTGCTGTAGCAGGGGGCTGTGATTATGGGTATTGCAGTGTACGCATA
TIGR4 TGGTACAATTATCTTTGCTGTAGCGGGGGCTGCGATTATGGGTATTGCAGTGTACGCATA
23FTW TGGTACAATTATCTTTGCTGTAGCGGGGGCTGTGATTATGGGTATTGCAGTGTACGCATA

Figure 196X

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14CSR TGTTAAAAACAACAAAGATGAGGATCAACTTGCTTAAGTAAGAGAGAAAAGGAGCCATTGA
670 TGTTAAAAACAACAAAGATGAGGATCAACTTGCTTAAGTAAGAGAGAAAAGGAGCCATTGA
6BF TGTTAAAAACAACAAAGATGAGGATCAACTTGCTTAAGTAAGAGAGAAAAGGAGCCATTGA
6BSP TGTTAAAAACAACAAAGATGAGGATCAACTTGCTTAAGTAAGAGAGAAAAGGAGCCATTGA
19AH TGTTAAAAACAACAAAGATGAGGATCAACTTGCTTAAGTAAGAGAGAAAAGGAGCCATTGA
23FPO TGTTAAAAACAACAAAGATGAGGATCAACTTGCTTAAGTAAGAGAGAAAAGGAGCCATTGA
19FTW TGTTAAAAACAACAAAGATGAGGATCAACTTGCTTAAGTAAGAGAGAAAAGGAGCCATTGA
9VSP TGTTAAAAACAACAAAGATGAGGATCAACTTGCTTAAGTAAGAGAGAAAAGGAGCCATTGA
TIGR4 TGTTAAAAACAACAAAGATGAGGATCAACTTGCTTAAGTAAGAGAGAAAAGGAGCCATTGA
23FTW TGTTAAAAACAACAAAGATGAGGATCAACTTGCTTAAGTAAGAGAGAAAAGGAGCCATTGA

14CSR TGACAATGCAGAAAATGCAGAAAATG-----
670 TGACAATGCAGAAAATGCAGAAAATG-----
6BF TGACAATGCAGAAAATGCAGAAAATG-----
6BSP TGACAATGCAGAAAATGCAGAAAATG-----
19AH TGACAATGCAGAAAATGCAGAAAATG-----
23FPO TGACAATGCAGAAAATGCAGAAAATG-----
19FTW TGACAATGCAGAAAATGCAGAAAATG-----
9VSP TGACAATGCAGAAAATGCAGAAAATGCAGAAAATGCAGAAAATGCAGAAAATGCAGAAAA
TIGR4 TGACAATGCAGAAAATGCAGAAAATG-----
23FTW TGACAATGCAGAAAATGCAGAAAATG-----

14CSR --ATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGATGGGGTGCACATG
670 --ATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGATGGGGTGCACATG
6BF --ATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGATGGGGTGCACATG
6BSP --ATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGATGGGGTGCACATG
19AH --ATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGATGGGGTGCACATG
23FPO --ATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGATGGGGTGCACATG
19FTW --ATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGATGGGGTGCACATG
9VSP TGATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGATGGGGTGCACATG
TIGR4 --ATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGATGGGGTGCACATG
23FTW --ATTAGTCGTATCTTCTTTGTTATGGCTCTGTGTTTTCTCTTGATGGGGTGCACATG

14CSR CAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAACATATCAGGAGGTGG
670 CAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAACATATCAGGAGGTGG
6BF CAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAACATATCAGGAGGTGG
6BSP CAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAACATATCAGGAGGTGG
19AH CAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAACATATCAGGAGGTGG
23FPO CAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAACATATCAGGAGGTGG
19FTW CAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAACATATCAGGAGGTGG
9VSP CAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAACATATCAGGAGGTGG
TIGR4 CAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAACATATCAGGAGGTGG
23FTW CAGTCCAAGCGCAAGAAGATCACACGTTGGTCTTGCAATTGGAGAACATATCAGGAGGTGG

14CSR TTAGTCAATTGCCATCTCGTGATGGTCATCGGTTGCAAGTATGGAAGTTGGATGATTTCGT
670 TTAGTCAATTGCCATCTCGTGATGGTCATCGGTTGCAAGTATGGAAGTTGGATGATTTCGT
6BF TTAGTCAATTGCCATCTCGTGATGGTCATCGGTTGCAAGTATGGAAGTTGGATGATTTCGT
6BSP TTAGTCAATTGCCATCTCGTGATGGTCATCGGTTGCAAGTATGGAAGTTGGATGATTTCGT
19AH TTAGTCAATTGCCATCTCGTGATGGTCATCGGTTGCAAGTATGGAAGTTGGATGATTTCGT
23FPO TTAGTCAATTGCCATCTCGTGATGGTCATCGGTTGCAAGTATGGAAGTTGGATGATTTCGT
19FTW TTAGTCAATTGCCATCTCGTGATGGTCATCGGTTGCAAGTATGGAAGTTGGATGATTTCGT
9VSP TTAGTCAATTGCCATCTCGTGATGGTCATCGGTTGCAAGTATGGAAGTTGGATGATTTCGT
TIGR4 TTAGTCAATTGCCATCTCGTGATGGTCATCGGTTGCAAGTATGGAAGTTGGATGATTTCGT
23FTW TTAGTCAATTGCCATCTCGTGATGGTCATCGGTTGCAAGTATGGAAGTTGGATGATTTCGT

Figure 196Y

14CSR ATTCCCTATGATGATCGGGTGCAAATTGTAAGAGACTTGCATTCTGTTGGGATGAGAATAAAC
670 ATTCCCTATGATGATCGGGTGCAAATTGTAAGAGACTTGCATTCTGTTGGGATGAGAATAAAC
6BF ATTCCCTATGATGATCGGGTGCAAATTGTAAGAGACTTGCATTCTGTTGGGATGAGAATAAAC
6BSP ATTCCCTATGATGATCGGGTGCAAATTGTAAGAGACTTGCATTCTGTTGGGATGAGAATAAAC
19AH ATTCCCTATGATGATCGGGTGCAAATTGTAAGAGACTTGCATTCTGTTGGGATGAGAATAAAC
23FPO ATTCCCTATGATAATCGGGTGCAAATTGTGAGAGACTTGCATTCTGTTGGGATGAGAATAAAC
19FTW ATTCCCTATGATAATCGGGTGCAAATTGTGAGAGACTTGCATTCTGTTGGGATGAGAATAAAC
9VSP ATTCCCTATGATAATCGGGTGCAAATTGTGAGAGACTTGCATTCTGTTGGGATGAGAATAAAC
TIGR4 ATTCCCTATGATGATCGGGTGCAAATTGTAAGAGACTTGCATTCTGTTGGGATGAGAATAAAC
23FTW ATTCCCTATGATAATCGGGTGCAAATTGTGAGAGACTTGCATTCTGTTGGGATGAGAATAAAC

14CSR TTTCTTCTTTCAAAAAGACTTCGTTTGAGATGACCTTCCTTGAGAATCAGATTGAAGTAT
670 TTTCTTCTTTCAAAAAGACTTCGTTTGAGATGACCTTCCTTGAGAATCAGATTGAAGTAT
6BF TTTCTTCTTTCAAAAAGACTTCGTTTGAGATGACCTTCCTTGAGAATCAGATTGAAGTAT
6BSP TTTCTTCTTTCAAAAAGACTTCGTTTGAGATGACCTTCCTTGAGAATCAGATTGAAGTAT
19AH TTTCTTCTTTCAAAAAGACTTCGTTTGAGATGACCTTCCTTGAGAATCAGATTGAAGTAT
23FPO TTTCTTCTTTCAAAAAGACTTCGTTTGAGATGACCTTCCTTGAGAATCAGATTGAAGTAT
19FTW TTTCTTCTTTCAAAAAGACTTCGTTTGAGATGACCTTCCTTGAGAATCAGATTGAAGTAT
9VSP TTTCTTCTTTCAAAAAGACTTCGTTTGAGATGACCTTCCTTGAGAATCAGATTGAAGTAT
TIGR4 TTTCTTCTTTCAAAAAGACTTCGTTTGAGATGACCTTCCTTGAGAATCAGATTGAAGTAT
23FTW TTTCTTCTTTCAAAAAGACTTCGTTTGAGATGACCTTCCTTGAGAATCAGATTGAAGTAT

14CSR CTCATATTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTT
670 CTCATATTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTT
6BF CTCATATTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTT
6BSP CTCATATTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTT
19AH CTCATATTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTT
23FPO CTCATATTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTT
19FTW CTCATATTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTT
9VSP CTCATATTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTT
TIGR4 CTCATATTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTT
23FTW CTCATATTCCAAATGGTCTTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTTCTT

14CSR ATCCAGCTGAATTTCTTTTTGAAATGACAGATCAAACGGTAGAGCCTTTGGTCATTGTAG
670 ATCCAGCTGAATTTCTTTTTGAAATGACAGATCAAACGGTAGAGCCTTTGGTCATTGTAG
6BF ATCCAGCTGAATTTCTTTTTGAAATGACAGATCAAACGGTAGAGCCTTTGGTCATTGTAG
6BSP ATCCAGCTGAATTTCTTTTTGAAATGACAGATCAAACGGTAGAGCCTTTGGTCATTGTAG
19AH ATCCAGCTGAATTTCTTTTTGAAATGACAGATCAAACGGTAGAGCCTTTGGTCATTGTAG
23FPO ATCCAGCTGAATTTCTTTTTGAAATGACAGATCAAACGGTAGAGCCTTTGGTCATTGTAG
19FTW ATCCAGCTGAATTTCTTTTTGAAATGACAGATCAAACGGTAGAGCCTTTGGTCATTGTAG
9VSP ATCCAGCTGAATTTCTTTTTGAAATGACAGATCAAACGGTAGAGCCTTTGGTCATTGTAG
TIGR4 ATCCAGCTGAATTTCTTTTTGAAATGACAGATCAAACGGTAGAGCCTTTGGTCATTGTAG
23FTW ATCCAGCTGAATTTCTTTTTGAAATGACAGATCAAACGGTAGAGCCTTTGGTCATTGTAG

14CSR CGAAAAAACAGATACAATGACAACAAAGGTGAAGCTGATAAAGGTGGATCAAGACCACA
670 CGAAAAAACAGATACAATGACAACAAAGGTGAAGCTGATAAAGGTGGATCAAGACCACA
6BF CGAAAAAACAGATACAATGACAACAAAGGTGAAGCTGATAAAGGTGGATCAAGACCACA
6BSP CGAAAAAACAGATACAATGACAACAAAGGTGAAGCTGATAAAGGTGGATCAAGACCACA
19AH CGAAAAAACAGATACAATGACAACAAAGGTGAAGCTGATAAAGGTGGATCAAGACCACA
23FPO CGAAAAAACAGATACGGTGACAACAAAGGTGAAGCTGATAAAGGTGGATCAAGACCACA
19FTW CGAAAAAACAGATACGGTGACAACAAAGGTGAAGCTGATAAAGGTGGATCAAGACCACA
9VSP CGAAAAAACAGATACGGTGACAACAAAGGTGAAGCTGATAAAGGTGGATCAAGACCACA
TIGR4 CGAAAAAACAGATACAATGACAACAAAGGTGAAGCTGATAAAGGTGGATCAAGACCACA
23FTW CGAAAAAACAGATACGGTGACAACAAAGGTGAAGCTGATAAAGGTGGATCAAGACCACA

Figure 196X

14CSR ATCGCTTGGAGGGTGTGGGCTTTAAATTGGTATCAGTAGCAAGAGATGGTTCTGAAAAAG
670 ATCGCTTGGAGGGTGTGGGCTTTAAATTGGTATCAGTAGCAAGAGATGGTTCTGAAAAAG
6BF ATCGCTTGGAGGGTGTGGGCTTTAAATTGGTATCAGTAGCAAGAGATGGTTCTGAAAAAG
6BSP ATCGCTTGGAGGGTGTGGGCTTTAAATTGGTATCAGTAGCAAGAGATGGTTCTGAAAAAG
19AH ATCGCTTGGAGGGTGTGGGCTTTAAATTGGTATCAGTAGCAAGAGATGGTTCTGAAAAAG
23FPO ATCGCTTGGAGGGTGTGGGCTTTAAATTGGTATCAGTAGCAAGAGATGGTTCTGAAAAAG
19FTW ATCGCTTGGAGGGTGTGGGCTTTAAATTGGTATCAGTAGCAAGAGATGGTTCTGAAAAAG
9VSP ATCGCTTGGAGGGTGTGGGCTTTAAATTGGTATCAGTAGCAAGAGATGGTTCTGAAAAAG
TIGR4 ATCGCTTGGAGGGTGTGGGCTTTAAATTGGTATCAGTAGCAAGAGATGGTTCTGAAAAAG
23FTW ATCGCTTGGAGGGTGTGGGCTTTAAATTGGTATCAGTAGCAAGAGATGGTTCTGAAAAAG

14CSR AGGTTCCCTTGATTGGAGAATACCGTTACAGTTCCTCTGGTCAAGTAGGGAGAACTCTCT
670 AGGTTCCCTTGATTGGAGAATACCGTTACAGTTCCTCTGGTCAAGTAGGGAGAACTCTCT
6BF AGGTTCCCTTGATTGGAGAATACCGTTACAGTTCCTCTGGTCAAGTAGGGAGAACTCTCT
6BSP AGGTTCCCTTGATTGGAGAATACCGTTACAGTTCCTCTGGTCAAGTAGGGAGAACTCTCT
19AH AGGTTCCCTTGATTGGAGAATACCGTTACAGTTCCTCTGGTCAAGTAGGGAGAACTCTCT
23FPO AGGTTCCCTTGATTGGAGAATACCGTTACAGTTCCTCTGGTCAAGTAGGGAGAACTCTCT
19FTW AGGTTCCCTTGATTGGAGAATACCGTTACAGTTCCTCTGGTCAAGTAGGGAGAACTCTCT
9VSP AGGTTCCCTTGATTGGAGAATACCGTTACAGTTCCTCTGGTCAAGTAGGGAGAACTCTCT
TIGR4 AGGTTCCCTTGATTGGAGAATACCGTTACAGTTCCTCTGGTCAAGTAGGGAGAACTCTCT
23FTW AGGTTCCCTTGATTGGAGAATACCGTTACAGTTCCTCTGGTCAAGTAGGGAGAACTCTCT

14CSR ATACTGATAAAAAATGGAGAGATTTTGTGACAAATCTTCCTCTTGGGAACATATCGTTTCA
670 ATACTGATAAAAAATGGAGAGATTTTGTGACAAATCTTCCTCTTGGGAACATATCGTTTCA
6BF ATACTGATAAAAAATGGAGAGATTTTGTGACAAATCTTCCTCTTGGGAACATATCGTTTCA
6BSP ATACTGATAAAAAATGGAGAGATTTTGTGACAAATCTTCCTCTTGGGAACATATCGTTTCA
19AH ATACTGATAAAAAATGGAGAGATTTTGTGACAAATCTTCCTCTTGGGAACATATCGTTTCA
23FPO ATACTGATAAAAAATGGAGAGATTTTGTGACAAATCTTCCTCTTGGGAACATATCGTTTCA
19FTW ATACTGATAAAAAATGGAGAGATTTTGTGACAAATCTTCCTCTTGGGAACATATCGTTTCA
9VSP ATACTGATAAAAAATGGAGAGATTTTGTGACAAATCTTCCTCTTGGGAACATATCGTTTCA
TIGR4 ATACTGATAAAAAATGGAGAGATTTTGTGACAAATCTTCCTCTTGGGAACATATCGTTTCA
23FTW ATACTGATAAAAAATGGAGAGATTTTGTGACAAATCTTCCTCTTGGGAACATATCGTTTCA

14CSR AGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGCTGG
670 AGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGCTGG
6BF AGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGCTGG
6BSP AGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGCTGG
19AH AGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGCTGG
23FPO AGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGTTGG
19FTW AGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGTTGG
9VSP AGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGTTGG
TIGR4 AGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGTTGG
23FTW AGGAGGTGGAGCCACTGGCAGGCTATGCTGTTACGACGCTGGATACGGATGTCCAGTTGG

14CSR TAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTG
670 TAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTG
6BF TAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTG
6BSP TAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTG
19AH TAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTG
23FPO TAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTG
19FTW TAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTG
9VSP TAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTG
TIGR4 TAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTG
23FTW TAGATCATCAGCTGGTGACGATTACGGTTGTCAATCAGAAATTACCACGTGGCAATGTTG

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14CSR ACTTTATGAAGGTGGATGGTCGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAGTCA
670 ACTTTATGAAGGTGGATGGTCGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAGTCA
6BF ACTTTATGAAGGTGGATGGTCGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAGTCA
6BSP ACTTTATGAAGGTGGATGGTCGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAGTCA
19AH ACTTTATGAAGGTGGATGGTCGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAGTCA
23FPO ACTTTATGAAGGTGGATGGTAGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAGTCA
19FTW ACTTTATGAAGGTGGATGGTAGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAGTCA
9VSP ACTTTATGAAGGTGGATGGTAGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAGTCA
TIGR4 ACTTTATGAAGGTGGATGGTCGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAGTCA
23FTW ACTTTATGAAGGTGGATGGTCGGACCAATACCTCTCTTCAAGGGGCAATGTTCAAAGTCA

14CSR TGAAAGAAGAAAGCGGACACTATACTCCTGTTCTTCAAAATGGTAAGGAAGTAGTTGTAA
670 TGAAAGAAGAAAGCGGACACTATACTCCTGTTCTTCAAAATGGTAAGGAAGTAGTTGTAA
6BF TGAAAGAAGAAAGCGGACACTATACTCCTGTTCTTCAAAATGGTAAGGAAGTAGTTGTAA
6BSP TGAAAGAAGAAAGCGGACACTATACTCCTGTTCTTCAAAATGGTAAGGAAGTAGTTGTAA
19AH TGAAAGAAGAAAGCGGACACTATACTCCTGTTCTTCAAAATGGTAAGGAAGTAGTTGTAA
23FPO TGAAAGAAGAAACGGGACACTATACTCCTGTTCTTCAAAATGGTAAGGAAGTAGTTGTGG
19FTW TGAAAGAAGAAACGGGACACTATACTCCTGTTCTTCAAAATGGTAAGGAAGTAGTTGTGG
9VSP TGAAAGAAGAAACGGGACACTATACTCCTGTTCTTCAAAATGGTAAGGAAGTAGTTGTGG
TIGR4 TGAAAGAAGAAAGCGGACACTATACTCCTGTTCTTCAAAATGGTAAGGAAGTAGTTGTAA
23FTW TGAAAGAAGAAACGGGACACTATACTCCTGTTCTTCAAAATGGTAAGGAAGTAGTTGTGG

14CSR CATCAGGGAAAGATGGTCGTTTCCGAGTGGAAGGTCTAGAGTATGGGACATACTATTTAT
670 CATCAGGGAAAGATGGTCGTTTCCGAGTGGAAGGTCTAGAGTATGGGACATACTATTTAT
6BF CATCAGGGAAAGATGGTCGTTTCCGAGTGGAAGGTCTAGAGTATGGGACATACTATTTAT
6BSP CATCAGGGAAAGATGGTCGTTTCCGAGTGGAAGGTCTAGAGTATGGGACATACTATTTAT
19AH CATCAGGGAAAGATGGTCGTTTCCGAGTGGAAGGTCTAGAGTATGGGACATACTATTTAT
23FPO CATCAGGGAAAGATGGTCGTTTCCGAGTGGAAGGTCTAGAGTATGGGACATACTATTTAT
19FTW CATCAGGGAAAGATGGTCGTTTCCGAGTGGAAGGTCTAGAGTATGGGACATACTATTTAT
9VSP CATCAGGGAAAGATGGTCGTTTCCGAGTGGAAGGTCTAGAGTATGGGACATACTATTTAT
TIGR4 CATCAGGGAAAGATGGTCGTTTCCGAGTGGAAGGTCTAGAGTATGGGACATACTATTTAT
23FTW CATCAGGGAAAGATGGTCGTTTCCGAGTGGAAGGTCTAGAGTATGGGACATACTATTTAT

14CSR GGGAGCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTTCCTTTACAATCG
670 GGGAGCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTTCCTTTACAATCG
6BF GGGAGCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTTCCTTTACAATCG
6BSP GGGAGCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTTCCTTTACAATCG
19AH GGGAGCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTTCCTTTACAATCG
23FPO GGGAGCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTTCCTTTACAATCG
19FTW GGGAGCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTTCCTTTACAATCG
9VSP GGGAGCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTTCCTTTACAATCG
TIGR4 GGGAGCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTTCCTTTACAATCG
23FTW GGGAGCTCCAAGCTCCAAGTGGTTATGTTCAATTAACATCGCCTGTTTCCTTTACAATCG

14CSR GGAAAGATACTCGTAAGGAACTGGTAACAGTGGTTAAAAATAACAAGCGACCACGGATTG
670 GGAAAGATACTCGTAAGGAACTGGTAACAGTGGTTAAAAATAACAAGCGACCACGGATTG
6BF GGAAAGATACTCGTAAGGAACTGGTAACAGTGGTTAAAAATAACAAGCGACCACGGATTG
6BSP GGAAAGATACTCGTAAGGAACTGGTAACAGTGGTTAAAAATAACAAGCGACCACGGATTG
19AH GGAAAGATACTCGTAAGGAACTGGTAACAGTGGTTAAAAATAACAAGCGACCACGGATTG
23FPO GGAAAGATACTCGTAAGGAACTGGTAACAGTGGTTAAAAATAACAAGCGACCACGGATTG
19FTW GGAAAGATACTCGTAAGGAACTGGTAACAGTGGTTAAAAATAACAAGCGACCACGGATTG
9VSP GGAAAGATACTCGTAAGGAACTGGTAACAGTGGTTAAAAATAACAAGCGACCACGGATTG
TIGR4 GGAAAGATACTCGTAAGGAACTGGTAACAGTGGTTAAAAATAACAAGCGACCACGGATTG
23FTW GGAAAGATACTCGTAAGGAACTGGTAACAGTGGTTAAAAATAACAAGCGACCACGGATTG

Figure 196AB

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14CSR ATGTGCCAGATACAGGGGAAGAAACCTTGTATATCTTGATGCTTGTGCCATTTTGTGT
670 ATGTGCCAGATACAGGGGAAGAAACCTTGTATATCTTGATGCTTGTGCCATTTTGTGT
6BF ATGTGCCAGATACAGGGGAAGAAACCTTGTATATCTTGATGCTTGTGCCATTTTGTGT
6BSP ATGTGCCAGATACAGGGGAAGAAACCTTGTATATCTTGATGCTTGTGCCATTTTGTGT
19AH ATGTGCCAGATACAGGGGAAGAAACCTTGTATATCTTGATGCTTGTGCCATTTTGTGT
23FPO ATGTGCCAGATACAGGGGAAGAAACCTTGTATATCTTGATGCTTGTGCCATTTTGTGT
19FTW ATGTGCCAGATACAGGGGAAGAAACCTTGTATATCTTGATGCTTGTGCCATTTTGTGT
9VSP ATGTGCCAGATACAGGGGAAGAAACCTTGTATATCTTGATGCTTGTGCCATTTTGTGT
TIGR4 ATGTGCCAGATACAGGGGAAGAAACCTTGTATATCTTGATGCTTGTGCCATTTTGTGT
23FTW ATGTGCCAGATACAGGGGAAGAAACCTTGTATATCTTGATGCTTGTGCCATTTTGTGT

14CSR TTGGTAGTGGTTATTATCTTACGAAAAAACCAATAACTGATATTCAATGTACATCATTA
670 TTGGTAGTGGTTATTATCTTACGAAAAAACCAATAACTGATATTCAATGTACATCATTA
6BF TTGGTAGTGGTTATTATCTTACGAAAAAACCAATAACTGATATTCAATGTACATCATTA
6BSP TTGGTAGTGGTTATTATCTTACGAAAAAACCAATAACTGATATTCAATGTACATCATTA
19AH TTGGTAGTGGTTATTATCTTACGAAAAAACCAATAACTGATATTCAATGTACATCATTA
23FPO TTGGTAGTGGCTATTATCTTACGAAAAAACCAATAACTGATATTCAATGTACATCATTA
19FTW TTGGTAGTGGCTATTATCTTACGAAAAAACCAATAACTGATATTCAATGTACATCATTA
9VSP TTGGTAGTGGCTATTATCTTACGAAAAAACCAATAACTGATATTCAATGTACATCATTA
TIGR4 TTGGTAGTGGCTATTATCTTACGAAAAAACCAATAACTGATATTCAATGTACATCATTA
23FTW TTGGTAGTGGCTATTATCTTACGAAAAAACCAATAACTGATATTCAATGTACATCATTA

14CSR TGAAAAAGATAGCAGGCTGAAGGGAAGACCAGAGTACTCTGAGGTGATGTTAATCAGGAA
670 TGAAAAAGATAGCAGGCTGAAGGGAAGACCAGAGTACTCTGAGGTGATGTTAATCAGGAA
6BF TGAAAAAGATAGCAGGCTGAAGGGAAGACCAGAGTACTCTGAGGTGATGTTAATCAGGAA
6BSP TGAAAAAGATAGCAGGCTGAAGGGAAGACCAGAGTACTCTGAGGTGATGTTAATCAGGAA
19AH TGAAAAAGATAGCAGGCTGAAGGGAAGACCAGAGTACTCTGAGGTGATGTTAATCAGGAA
23FPO TGAAAAAGATAGCAGGCTGAAGGGAAGACCAGAGTACTCTGAGGTGATGTTAATCAGGAA
19FTW TGAAAAAGATAGCAGGCTGAAGGGAAGACCAGAGTACTCTGAGGTGATGTTAATCAGGAA
9VSP TGAAAAAGATAGCAGGCTGAAGGGAAGACCAGAGTACTCTGAGGTGATGTTAATCAGGAA
TIGR4 TGAAAAAGATAGCAGGCTGAAGGGAAGACCAGAGTACTCTGAGGTGATGTTAATCAGGAA
23FTW TGAAAAAGATAGCAGGCTGAAGGGAAGACCAGAGTACTCTGAGGTGATGTTAATCAGGAA

14CSR TCATGGTGATGTGGCATGAATCACAATAACGGATATGAGGCTGGGCAGATTGTGCCAGCC
670 TCATGGTGATGTGGCATGAATCACAATAACGGATATGAGGCTGGGCAGATTGTGCCAGCC
6BF TCATGGTGATGTGGCATGAATCACAATAACGGATATGAGGCTGGGCAGATTGTGCCAGCC
6BSP TCATGGTGATGTGGCATGAATCACAATAACGGATATGAGGCTGGGCAGATTGTGCCAGCC
19AH TCATGGTGATGTGGCATGAATCACAATAACGGATATGAGGCTGGGCAGATTGTGCCAGCC
23FPO TCATGGTGATTGTGGCATGAATCATAATAACGGATATGAGGCTGGGCAGATTGTGCCAGCC
19FTW TCATGGTGATTGTGGCATGAATCATAATAACGGATATGAGGCTGGGCAGATTGTGCCAGCC
9VSP TCATGGTGATTGTGGCATGAATCATAATAACGGATATGAGGCTGGGCAGATTGTGCCAGCC
TIGR4 TCATGGTGATGTGGCATGAATCACAATAACGGATATGAGGCTGGGCAGATTGTGCCAGCC
23FTW TCATGGTGATTGTGGCATGAATCACAATAACGGATATGAGGCTGGGCAGATTGTGCCAGCC

14CSR TCATTGTGGGTTATTGTTTGTAAAACGATAGGACTGGTCTGGTAATCATTTTAGGAATGG
670 TCATTGTGGGTTATTGTTTGTAAAACGATAGGACTGGTCTGGTAATCATTTTAGGAATGG
6BF TCATTGTGGGTTATTGTTTGTAAAACGATAGGACTGGTCTGGTAATCATTTTAGGAATGG
6BSP TCATTGTGGGTTATTGTTTGTAAAACGATAGGACTGGTCTGGTAATCATTTTAGGAATGG
19AH TCATTGTGGGTTATTGTTTGTAAAACGATAGGACTGGTCTGGTAATCATTTTAGGAATGG
23FPO TCATTGTGGGTTATTGTTTGTAAAACGATAGGACTGGTCTGGTAATCATTTTAGGAATGG
19FTW TCATTGTGGGTTATTGTTTGTAAAACGATAGGACTGGTCTGGTAATCATTTTAGGAATGG
9VSP TCATTGTGGGTTATTGTTTGTAAAACGATAGGACTGGTCTGGTAATCATTTTAGGAATGG
TIGR4 TCATTGTGGGTTATTGTTTGTAAAACGATAGGACTGGTCTGGTAATCATTTTAGGAATGG
23FTW TCATTGTGGGTTATTGTTTGTAAAACGATAGGACTGGTCTGGTAATCATTTTAGGAATGG

Figure 196AC

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14CSR ACAGGACTGGGATTCTGATTTAAATGGATGGTGAATCAGAAAGAAATGAGATTTTCTCG
670 ACAGGACTGGGATTCTGATTTAAATGGATGGTGAATCAGAAAGAAATGAGATTTTCTCG
6BF ACAGGACTGGGATTCTGATTTAAATGGATGGTGAATCAGAAAGAAATGAGATTTTCTCG
6BSP ACAGGACTGGGATTCTGATTTAAATGGATGGTGAATCAGAAAGAAATGAGATTTTCTCG
19AH ACAGGACTGGGATTCTGATTTAAATGGATGGTGAATCAGAAAGAAATGAGATTTTCTCG
23FPO ACAGGACTGGGATTCTGATTTAAATGGATGGTGAATCAGAAAGAAATGAGATTTTCTCG
19FTW ACAGGACTGGGATTCTGATTTAAATGGATGGTGAATCAGAAAGAAATGAGATTTTCTCG
9VSP ACAGGACTGGGATTCTGATTTAAATGGATGGTGAATCAGAAAGAAATGAGATTTTCTCG
TIGR4 ACAGGACTGGGATTCTGATTTAAATGGATGGTGAATCAGAAAGAAATGAGATTTTCTCG
23FTW ACAGGACTGGGATTCTGATTTAAATGGATGGTGAATCAGAAAGAAATGAGATTTTCTCG

14CSR TTTCTCTTAGCAGATAGGATTGTCTGTTAGGAAAAGCGATAAAATGATGAGTTTGAAGAT
670 TTTCTCTTAGCAGATAGGATTGTCTGTTAGGAAAAGCGATAAAATGATGAGTTTGAAGAT
6BF TTTCTCTTAGCAGATAGGATTGTCTGTTAGGAAAAGCGATAAAATGATGAGTTTGAAGAT
6BSP TTTCTCTTAGCAGATAGGATTGTCTGTTAGGAAAAGCGATAAAATGATGAGTTTGAAGAT
19AH TTTCTCTTAGCAGATAGGATTGTCTGTTAGGAAAAGCGATAAAATGATGAGTTTGAAGAT
23FPO TTTCTCTTAGCAGATAGGATTGTCTGTTAGGAAAAGCGATAAAATGATGAGTTTGAAGAT
19FTW TTTCTCTTAGCAGATAGGATTGTCTGTTAGGAAAAGCGATAAAATGATGAGTTTGAAGAT
9VSP TTTCTCTTAGCAGATAGGATTGTCTGTTAGGAAAAGCGATAAAATGATGAGTTTGAAGAT
TIGR4 TTTCTCTTAGCAGATAGGATTGTCTGTTAGGAAAAGCGATAAAATGATGAGTTTGAAGAT
23FTW TTTCTCTTAGCAGATAGGATTGTCTGTTAGGAAAAGCGATAAAATGATGAGTTTGAAGAT

14CSR AAAGGGATGCTGATAAAAA-TGGTAAAAACAAAAAGCAAAAACGAAATAATCTCCTATT
670 AAAGGGATGCTGATAAAAA-TGGTAAAAACAAAAAGCAAAAACGAAATAATCTCCTATT
6BF AAAGGGATGCTGATAAAAA-TGGTAAAAACAAAAAGCAAAAACGAAATAATCTCCTATT
6BSP AAAGGGATGCTGATAAAAA-TGGTAAAAACAAAAAGCAAAAACGAAATAATCTCCTATT
19AH AAAGGGATGCTGATAAAAA-TGGTAAAAACAAAAAGCAAAAACGAAATAATCTCCTATT
23FPO AAAGGAATGCTGATAAAAAATGGCAAAAACAAAAAGCAAAAACGAAACAATCTCCTATT
19FTW AAAGGAATGCTGATAAAAAATGGCAAAAACAAAAAGCAAAAACGAAACAATCTCCTATT
9VSP AAAGGAATGCTGATAAAAAATGGCAAAAACAAAAAGCAAAAACGAAACAATCTCCTATT
TIGR4 AAAGGGATGCTGATAAAAA-TGGTAAAAACAAAAAGCAAAAACGAAATAATCTCCTATT
23FTW AAAGGGATGCTGATAAAAA-TGGTAAAAACAAAAAGCAAAAACGAAATAATCTCCTATT

14CSR AGGAGTGGTATTTTTTCATTGGAATGGCGGTAATGGCGTATCCGCTGGTGTCTCGCTTGTA
670 AGGAGTGGTATTTTTTCATTGGAATGGCGGTAATGGCGTATCCGCTGGTGTCTCGCTTGTA
6BF AGGAGTGGTATTTTTTCATTGGAATGGCGGTAATGGCGTATCCGCTGGTGTCTCGCTTGTA
6BSP AGGAGTGGTATTTTTTCATTGGAATGGCGGTAATGGCGTATCCGCTGGTGTCTCGCTTGTA
19AH AGGAGTGGTATTTTTTCATTGGAATGGCGGTAATGGCGTATCCGCTGGTGTCTCGCTTGTA
23FPO AGGAGTGGTATTTTTTCATTGGAATGGCGGTAATGGCGTATCCGCTGGTGTCTCGCTTGTA
19FTW AGGAGTGGTATTTTTTCATTGGAATGGCGGTAATGGCGTATCCGCTGGTGTCTCGCTTGTA
9VSP AGGAGTGGTATTTTTTCATTGGAATGGCGGTAATGGCGTATCCGCTGGTGTCTCGCTTGTA
TIGR4 AGGAGTGGTATTTTTTCATTGGAATGGCGGTAATGGCGTATCCGCTGGTGTCTCGCTTGTA
23FTW AGGAGTGGTATTTTTTCATTGGAATGGCGGTAATGGCGTATCCGCTGGTGTCTCGCTTGTA

14CSR TTATCGAGTGAATCAAATCAACAAATTGCTGACTTTGATAAGGAAAAAGCAACGTTGGA
670 TTATCGAGTGAATCAAATCAACAAATTGCTGACTTTGATAAGGAAAAAGCAACGTTGGA
6BF TTATCGAGTGAATCAAATCAACAAATTGCTGACTTTGATAAGGAAAAAGCAACGTTGGA
6BSP TTATCGAGTGAATCAAATCAACAAATTGCTGACTTTGATAAGGAAAAAGCAACGTTGGA
19AH TTATCGAGTGAATCAAATCAACAAATTGCTGACTTTGATAAGGAAAAAGCAACGTTGGA
23FPO TTATCGAGTGAATCAAATCAACAAATTGCTGACTTTGATAAGGAAAAAGCAACGTTGGA
19FTW TTATCGAGTGAATCAAATCAACAAATTGCTGACTTTGATAAGGAAAAAGCAACGTTGGA
9VSP TTATCGAGTGAATCAAATCAACAAATTGCTGACTTTGATAAGGAAAAAGCAACGTTGGA
TIGR4 TTATCGAGTGAATCAAATCAACAAATTGCTGACTTTGATAAGGAAAAAGCAACGTTGGA
23FTW TTATCGAGTGAATCAAATCAACAAATTGCTGACTTTGATAAGGAAAAAGCAACGTTGGA

Figure 196AD

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14CSR TGAGGCTGACATTGATGAACGAATGAAATTGGCACAAGCCTTCAATGACTCTTTGAATAA
670 TGAGGCTGACATTGATGAACGAATGAAATTGGCACAAGCCTTCAATGACTCTTTGAATAA
6BF TGAGGCTGACATTGATGAACGAATGAAATTGGCACAAGCCTTCAATGACTCTTTGAATAA
6BSP TGAGGCTGACATTGATGAACGAATGAAATTGGCACAAGCCTTCAATGACTCTTTGAATAA
19AH TGAGGCTGACATTGATGAACGAATGAAATTGGCACAAGCCTTCAATGACTCTTTGAATAA
23FPO TGAGGCTGACATTGATGAACGAATGAAATTGGCACAAGCCTTCAATGACTCTTTGAATAA
19FTW TGAGGCTGACATTGATGAACGAATGAAATTGGCACAAGCCTTCAATGACTCTTTGAATAA
9VSP TGAGGCTGACATTGATGAACGAATGAAATTGGCACAAGCCTTCAATGACTCTTTGAATAA
TIGR4 TGAGGCTGACATTGATGAACGAATGAAATTGGCACAAGCCTTCAATGACTCTTTGAATAA
23FTW TGAGGCTGACATTGATGAACGAATGAAATTGGCACAAGCCTTCAATGACTCTTTGAATAA

14CSR TGTAGTGAGTGGCGATCCTTGGTCGGAAGAAATGAAGAAAAAGGGCGAGCAGAGTATGC
670 TGTAGTGAGTGGCGATCCTTGGTCGGAAGAAATGAAGAAAAAGGGCGAGCAGAGTATGC
6BF TGTAGTGAGTGGCGATCCTTGGTCGGAAGAAATGAAGAAAAAGGGCGAGCAGAGTATGC
6BSP TGTAGTGAGTGGCGATCCTTGGTCGGAAGAAATGAAGAAAAAGGGCGAGCAGAGTATGC
19AH TGTAGTGAGTGGCGATCCTTGGTCGGAAGAAATGAAGAAAAAGGGCGAGCAGAGTATGC
23FPO TGTAGTGAGTGGCGATCCTTGGTCGGAAGAAATGAAGAAAAAGGGCGAGCAGAGTATGC
19FTW TGTAGTGAGTGGCGATCCTTGGTCGGAAGAAATGAAGAAAAAGGGCGAGCAGAGTATGC
9VSP TGTAGTGAGTGGCGATCCTTGGTCGGAAGAAATGAAGAAAAAGGGCGAGCAGAGTATGC
TIGR4 TGTAGTGAGTGGCGATCCTTGGTCGGAAGAAATGAAGAAAAAGGGCGAGCAGAGTATGC
23FTW TGTAGTGAGTGGCGATCCTTGGTCGGAAGAAATGAAGAAAAAGGGCGAGCAGAGTATGC

14CSR ACGTATGTTAGAAATCCATGAGCGGATGGGGCATGTGGAAATCCCCGTTATTGACGTGGA
670 ACGTATGTTAGAAATCCATGAGCGGATGGGGCATGTGGAAATCCCCGTTATTGACGTGGA
6BF ACGTATGTTAGAAATCCATGAGCGGATGGGGCATGTGGAAATCCCCGTTATTGACGTGGA
6BSP ACGTATGTTAGAAATCCATGAGCGGATGGGGCATGTGGAAATCCCCGTTATTGACGTGGA
19AH ACGTATGTTAGAAATCCATGAGCGGATGGGGCATGTGGAAATCCCCGTTATTGACGTGGA
23FPO ACGTATGTTAGAAATCCATGAGCGGATGGGGCATGTGGAAATCCCCGTTATTGACGTGGA
19FTW ACGTATGTTAGAAATCCATGAGCGGATGGGGCATGTGGAAATCCCCGTTATTGACGTGGA
9VSP ACGTATGTTAGAAATCCATGAGCGGATGGGGCATGTGGAAATCCCCGTTATTGACGTGGA
TIGR4 ACGTATGTTAGAAATCCATGAGCGGATGGGGCATGTGGAAATCCCCGTTATTGACGTGGA
23FTW ACGTATGTTAGAAATCCATGAGCGGATGGGGCATGTGGAAATCCCCGTTATTGACGTGGA
*** ***** *

14CSR TTTGCCGGTTTATGCTGGTACTGCTGAAGAGGTATTGCAGCAAGGGGCTGGGCATCTAGA
670 TTTGCCGGTTTATGCTGGTACTGCTGAAGAGGTATTGCAGCAAGGGGCTGGGCATCTAGA
6BF TTTGCCGGTTTATGCTGGTACTGCTGAAGAGGTATTGCAGCAAGGGGCTGGGCATCTAGA
6BSP TTTGCCGGTTTATGCTGGTACTGCTGAAGAGGTATTGCAGCAAGGGGCTGGGCATCTAGA
19AH TTTGCCGGTTTATGCTGGTACTGCTGAAGAGGTATTGCAGCAAGGGGCTGGGCATCTAGA
23FPO TTTGCCGGTTTATGCTGGTACTGCTGAAGAGGTATTGCAGCAAGGGGCTGGGCATCTAGA
19FTW TTTGCCGGTTTATGCTGGTACTGCTGAAGAGGTATTGCAGCAAGGGGCTGGGCATCTAGA
9VSP TTTGCCGGTTTATGCTGGTACTGCTGAAGAGGTATTGCAGCAAGGGGCTGGGCATCTAGA
TIGR4 TTTGCCGGTTTATGCTGGTACTGCTGAAGAGGTATTGCAGCAAGGGGCTGGGCATCTAGA
23FTW TTTGCCGGTTTATGCTGGTACTGCTGAAGAGGTATTGCAGCAAGGGGCTGGGCATCTAGA

14CSR GGGAACTTCTCTGCCGATCGGAGGCAATTCGACCCATGCGGTGATTACGGCACATACAGG
670 GGGAACTTCTCTGCCGATCGGAGGCAATTCGACCCATGCGGTGATTACGGCACATACAGG
6BF GGGAACTTCTCTGCCGATCGGAGGCAATTCGACCCATGCGGTGATTACGGCACATACAGG
6BSP GGGAACTTCTCTGCCGATCGGAGGCAATTCGACCCATGCGGTGATTACGGCACATACAGG
19AH GGGAACTTCTCTGCCGATCGGAGGCAATTCGACCCATGCGGTGATTACGGCACATACAGG
23FPO GGGAACTTCTCTACCGATCGGAGGCAATTCGACCCATGCGGTGATTACGGCACATACAGG
19FTW GGGAACTTCTCTACCGATCGGAGGCAATTCGACCCATGCGGTGATTACGGCACATACAGG
9VSP GGGAACTTCTCTACCGATCGGAGGCAATTCGACCCATGCGGTGATTACGGCACATACAGG
TIGR4 GGGAACTTCTCTGCCGATCGGAGGCAATTCGACCCATGCGGTGATTACGGCACATACAGG
23FTW GGGAACTTCTCTGCCGATCGGAGGCAATTCGACCCATGCGGTGATTACGGCACATACAGG

Figure 196AE

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14CSR TTTGCCAACAGCTAAGATGTTTACGGATTGACCAAACCTTAAAGTTGGGGATAAGTTTTA
670 TTTGCCAACAGCTAAGATGTTTACGGATTGACCAAACCTTAAAGTTGGGGATAAGTTTTA
6BF TTTGCCAACAGCTAAGATGTTTACGGATTGACCAAACCTTAAAGTTGGGGATAAGTTTTA
6BSP TTTGCCAACAGCTAAGATGTTTACGGATTGACCAAACCTTAAAGTTGGGGATAAGTTTTA
19AH TTTGCCAACAGCTAAGATGTTTACGGATTGACCAAACCTTAAAGTTGGGGATAAGTTTTA
23FPO TTTGCCAACGGCTAAGATGTTTACGGATTGACCAAACCTTAAAGTTGGGGATAAGTTTTA
19FTW TTTGCCAACGGCTAAGATGTTTACGGATTGACCAAACCTTAAAGTTGGGGATAAGTTTTA
9VSP TTTGCCAACGGCTAAGATGTTTACGGATTGACCAAACCTTAAAGTTGGGGATAAGTTTTA
TIGR4 TTTGCCAACAGCTAAGATGTTTACGGATTGACCAAACCTTAAAGTTGGGGATAAGTTTTA
23FTW TTTGCCAACAGCTAAGATGTTTACGGATTGACCAAACCTTAAAGTTGGGGATAAGTTTTA
***** **

14CSR TGTGCACAATATCAAGGAAGTGATGGCCTATCAAGTGGATCAAGTAAAGGTGATTGAGCC
670 TGTGCACAATATCAAGGAAGTGATGGCCTATCAAGTGGATCAAGTAAAGGTGATTGAGCC
6BF TGTGCACAATATCAAGGAAGTGATGGCCTATCAAGTGGATCAAGTAAAGGTGATTGAGCC
6BSP TGTGCACAATATCAAGGAAGTGATGGCCTATCAAGTGGATCAAGTAAAGGTGATTGAGCC
19AH TGTGCACAATATCAAGGAAGTGATGGCCTATCAAGTGGATCAAGTAAAGGTGATTGAGCC
23FPO TGTGCACAATATCAAGGAAGTGATGGCCTATCAAGTGGATCAAGTAAAGGTGATTGAGCC
19FTW TGTGCACAATATCAAGGAAGTGATGGCCTATCAAGTGGATCAAGTAAAGGTGATTGAGCC
9VSP TGTGCACAATATCAAGGAAGTGATGGCCTATCAAGTGGATCAAGTAAAGGTGATTGAGCC
TIGR4 TGTGCACAATATCAAGGAAGTGATGGCCTATCAAGTGGATCAAGTAAAGGTGATTGAGCC
23FTW TGTGCACAATATCAAGGAAGTGATGGCCTATCAAGTGGATCAAGTAAAGGTGATTGAGCC

14CSR GACGAACTTTGATGATTTATTGATTGTACCAGGTCATGATTATGTGACCTTGCTGACTTG
670 GACGAACTTTGATGATTTATTGATTGTACCAGGTCATGATTATGTGACCTTGCTGACTTG
6BF GACGAACTTTGATGATTTATTGATTGTACCAGGTCATGATTATGTGACCTTGCTGACTTG
6BSP GACGAACTTTGATGATTTATTGATTGTACCAGGTCATGATTATGTGACCTTGCTGACTTG
19AH GACGAACTTTGATGATTTATTGATTGTACCAGGTCATGATTATGTGACCTTGCTGACTTG
23FPO GACGAACTTTGATGATTTATTGATTGTACCAGGTCATGATTATGTGACCTTGCTGACTTG
19FTW GACGAACTTTGATGATTTATTGATTGTACCAGGTCATGATTATGTGACCTTGCTGACTTG
9VSP GACGAACTTTGATGATTTATTGATTGTACCAGGTCATGATTATGTGACCTTGCTGACTTG
TIGR4 GACGAACTTTGATGATTTATTGATTGTACCAGGTCATGATTATGTGACCTTGCTGACTTG
23FTW GACGAACTTTGATGATTTATTGATTGTACCAGGTCATGATTATGTGACCTTGCTGACTTG
***** **

14CSR TACGCCATACATGATCAATACCCATCGTCTATTGGTTCGGGGGCATCGGATACCGTACGT
670 TACGCCATACATGATCAATACCCATCGTCTATTGGTTCGGGGGCATCGGATACCGTACGT
6BF TACGCCATACATGATCAATACCCATCGTCTATTGGTTCGGGGGCATCGGATACCGTACGT
6BSP TACGCCATACATGATCAATACCCATCGTCTATTGGTTCGGGGGCATCGGATACCGTACGT
19AH TACGCCATACATGATCAATACCCATCGTCTATTGGTTCGGGGGCATCGGATACCGTACGT
23FPO TACGCCATACATGATCAATACCCATCGTCTATTGGTTCGGGGGCATCGGATACCGTACGT
19FTW TACGCCATACATGATCAATACCCATCGTCTATTGGTTCGGGGGCATCGGATACCGTACGT
9VSP TACGCCATACATGATCAATACCCATCGTCTATTGGTTCGGGGGCATCGGATACCGTACGT
TIGR4 TACGCCATACATGATCAATACCCATCGTCTATTGGTTCGGGGGCATCGGATACCGTACGT
23FTW TACGCCATACATGATCAATACCCATCGTCTATTGGTTCGGGGGCATCGGATACCGTACGT

14CSR AGCAGAGGTTGAGGAAGAATTTATTGCAGCAAACAACTCAGTCATCTCTATCGCTACCT
670 AGCAGAGGTTGAGGAAGAATTTATTGCAGCAAACAACTCAGTCATCTCTATCGCTACCT
6BF AGCAGAGGTTGAGGAAGAATTTATTGCAGCAAACAACTCAGTCATCTCTATCGCTACCT
6BSP AGCAGAGGTTGAGGAAGAATTTATTGCAGCAAACAACTCAGTCATCTCTATCGCTACCT
19AH AGCAGAGGTTGAGGAAGAATTTATTGCAGCAAACAACTCAGTCATCTCTATCGCTACCT
23FPO AGCAGAGGTTGAGGAAGAATTTATTGCGGCAAACAACTCAGTCATCTCTATCGCTACCT
19FTW AGCAGAGGTTGAGGAAGAATTTATTGCGGCAAACAACTCAGTCATCTCTATCGCTACCT
9VSP AGCAGAGGTTGAGGAAGAATTTATTGCGGCAAACAACTCAGTCATCTCTATCGCTACCT
TIGR4 AGCAGAGGTTGAGGAAGAATTTATTGCAGCAAACAACTCAGTCATCTCTATCGCTACCT
23FTW AGCAGAGGTTGAGGAAGAATTTATTGCAGCAAACAACTCAGTCATCTCTATCGCTACCT

Figure 196AF

14CSR GTTTTATGTGGCAGTTGGTTTGATTGTGATTCTTTTATGGATTATTTCGACGCTTGCGCAA
670 GTTTTATGTGGCAGTTGGTTTGATTGTGATTCTTTTATGGATTATTTCGACGCTTGCGCAA
6BF GTTTTATGTGGCAGTTGGTTTGATTGTGATTCTTTTATGGATTATTTCGACGCTTGCGCAA
6BSP GTTTTATGTGGCAGTTGGTTTGATTGTGATTCTTTTATGGATTATTTCGACGCTTGCGCAA
19AH GTTTTATGTGGCAGTTGGTTTGATTGTGATTCTTTTATGGATTATTTCGACGCTTGCGCAA
23FPO GTTTTATGTGGCAGTTGGTTTGATTGTGATTCTTTTATGGATTATTTCGACGCTTGCGCAA
19FTW GTTTTATGTGGCAGTTGGTTTGATTGTGATTCTTTTATGGATTATTTCGACGCTTGCGCAA
9VSP GTTTTATGTGGCAGTTGGTTTGATTGTGATTCTTTTATGGATTATTTCGACGCTTGCGCAA
TIGR4 GTTTTATGTGGCAGTTGGTTTGATTGTGATTCTTTTATGGATTATTTCGACGCTTGCGCAA
23FTW GTTTTATGTGGCAGTTGGTTTGATTGTGATTCTTTTATGGATTATTTCGACGCTTGCGCAA

14CSR GAAGAAAAACAACCGGAAAAGGCTTTGAAGGCGCTGAAAGCAGCAAGGAAGGAAGTGAA
670 GAAGAAAAACAACCGGAAAAGGCTTTGAAGGCGCTGAAAGCAGCAAGGAAGGAAGTGAA
6BF GAAGAAAAACAACCGGAAAAGGCTTTGAAGGCGCTGAAAGCAGCAAGGAAGGAAGTGAA
6BSP GAAGAAAAACAACCGGAAAAGGCTTTGAAGGCGCTGAAAGCAGCAAGGAAGGAAGTGAA
19AH GAAGAAAAACAACCGGAAAAGGCTTTGAAGGCGCTGAAAGCAGCAAGGAAGGAAGTGAA
23FPO GAAGAAACGGCAATCAGAAAGAGCTTTGAAAGCATTGAAGGAAGCTACTAAGGAAGTGAA
19FTW GAAGAAACGGCAATCAGAAAGAGCTTTGAAAGCATTGAAGGAAGCTACTAAGGAAGTGAA
9VSP GAAGAAACGGCAATCAGAAAGAGCTTTGAAAGCATTGAAGGAAGCTACTAAGGAAGTGAA
TIGR4 GAAGAAAAACAACCGGAAAAGGCTTTGAAGGCGCTGAAAGCAGCAAGGAAGGAAGTGAA
23FTW GAAGAAAAACAACCGGAAAAGGCTTTGAAGGCGCTGAAAGCAGCAAGGAAGGAAGTGAA
***** * * * * *

14CSR GGTGGAGGATGGACAACAGTAGACGTTTACGAAAAAAGGCACAAAAAAGAAGAAACATC
670 GGTGGAGGATGGACAACAGTAGACGTTTACGAAAAAAGGCACAAAAAAGAAGAAACATC
6BF GGTGGAGGATGGACAACAGTAGACGTTTACGAAAAAAGGCACAAAAAAGAAGAAACATC
6BSP GGTGGAGGATGGACAACAGTAGACGTTTACGAAAAAAGGCACAAAAAAGAAGAAACATC
19AH GGTGGAGGATGGACAACAGTAGACGTTTACGAAAAAAGGCACAAAAAAGAAGAAACATC
23FPO GGTAGAGGATGAGTAAGAGTAGATATTACGGAAGAAAGAGCGTGAAAAAGAAGAAATC
19FTW GGTAGAGGATGAGTAAGAGTAGATATTACGGAAGAAAGAGCGTGAAAAAGAAGAAATC
9VSP GGTAGAGGATGAGTAAGAGTAGATATTACGGAAGAAAGAGCGTGAAAAAGAAGAAATC
TIGR4 GGTGGAGGATGGACAACAGTAGACGTTTACGAAAAAAGGCACAAAAAAGAAGAAACATC
23FTW GGTGGAGGATGGACAACAGTAGACGTTTACGAAAAAAGGCACAAAAAAGAAGAAACATC
*** * * * * *

14CSR CGCTGATCCTTCTTCTGATTTTCTTAGTAGGATTGCGCGTTGCGATATATCCATTGGTGT
670 CGCTGATCCTTCTTCTGATTTTCTTAGTAGGATTGCGCGTTGCGATATATCCATTGGTGT
6BF CGCTGATCCTTCTTCTGATTTTCTTAGTAGGATTGCGCGTTGCGATATATCCATTGGTGT
6BSP CGCTGATCCTTCTTCTGATTTTCTTAGTAGGATTGCGCGTTGCGATATATCCATTGGTGT
19AH CGCTGATCCTTCTTCTGATTTTCTTAGTAGGATTGCGCGTTGCGATATATCCATTGGTGT
23FPO CGTTCATTCTTCTTCTGATTTTTTGGTGGGGCTTGCCGTTGCGATGTATCCCTTGGTGT
19FTW CGTTCATTCTTCTTCTGATTTTTTGGTGGGGCTTGCCGTTGCGATGTATCCCTTGGTGT
9VSP CGTTCATTCTTCTTCTGATTTTTTGGTGGGGCTTGCCGTTGCGATGTATCCCTTGGTGT
TIGR4 CGCTGATCCTTCTTCTGATTTTCTTAGTAGGATTGCGCGTTGCGATATATCCATTGGTGT
23FTW CGCTGATCCTTCTTCTGATTTTCTTAGTAGGATTGCGCGTTGCGATATATCCATTGGTGT
* * * * *

14CSR CTCGTTATTATTATCGTATTGAGTCAAACGAGGTTATTAAGAGTTTGATGAGACGGTTT
670 CTCGTTATTATTATCGTATTGAGTCAAACGAGGTTATTAAGAGTTTGATGAGACGGTTT
6BF CTCGTTATTATTATCGTATTGAGTCAAACGAGGTTATTAAGAGTTTGATGAGACGGTTT
6BSP CTCGTTATTATTATCGTATTGAGTCAAACGAGGTTATTAAGAGTTTGATGAGACGGTTT
19AH CTCGTTATTATTATCGTATTGAGTCAAACGAGGTTATTAAGAGTTTGATGAGACGGTTT
23FPO CTCGTTATTATTATCGTATTGAGTCAAACGAGGTTATTAAGAGTTTGATGAGACGGTTT
19FTW CTCGTTATTATTATCGTATTGAGTCAAACGAGGTTATTAAGAGTTTGATGAGACGGTTT
9VSP CTCGTTATTATTATCGTATTGAGTCAAACGAGGTTATTAAGAGTTTGATGAGACGGTTT
TIGR4 CTCGTTATTATTATCGTATTGAGTCAAACGAGGTTATTAAGAGTTTGATGAGACGGTTT
23FTW CTCGTTATTATTATCGTATTGAGTCAAACGAGGTTATTAAGAGTTTGATGAGACGGTTT

14CSR CCCAGATGGATAAGGCAGAACTTGAGGAGCGTTGGCGCTTGGCTCAAGCCTTCAATGCGA
670 CCCAGATGGATAAGGCAGAACTTGAGGAGCGTTGGCGCTTGGCTCAAGCCTTCAATGCGA
6BF CCCAGATGGATAAGGCAGAACTTGAGGAGCGTTGGCGCTTGGCTCAAGCCTTCAATGCGA
6BSP CCCAGATGGATAAGGCAGAACTTGAGGAGCGTTGGCGCTTGGCTCAAGCCTTCAATGCGA
19AH CCCAGATGGATAAGGCAGAACTTGAGGAGCGTTGGCGCTTGGCTCAAGCCTTCAATGCGA
23FPO CCCAGATGGATAAGGCAGAACTTGAGGAGCGTTGGCGCTTGGCTCAAGCCTTCAATGCGA
19FTW CCCAGATGGATAAGGCAGAACTTGAGGAGCGTTGGCGCTTGGCTCAAGCCTTCAATGCGA
9VSP CCCAGATGGATAAGGCAGAACTTGAGGAGCGTTGGCGCTTGGCTCAAGCCTTCAATGCGA
TIGR4 CCCAGATGGATAAGGCAGAACTTGAGGAGCGTTGGCGCTTGGCTCAAGCCTTCAATGCGA
23FTW CCCAGATGGATAAGGCAGAACTTGAGGAGCGTTGGCGCTTGGCTCAAGCCTTCAATGCGA

14CSR CCTTGAAACCATCTGAAATTCCTGATCCTTTTACAGAGCAAGAGAAAAAGAAAGGCGTCT
670 CCTTGAAACCATCTGAAATTCCTGATCCTTTTACAGAGCAAGAGAAAAAGAAAGGCGTCT
6BF CCTTGAAACCATCTGAAATTCCTGATCCTTTTACAGAGCAAGAGAAAAAGAAAGGCGTCT
6BSP CCTTGAAACCATCTGAAATTCCTGATCCTTTTACAGAGCAAGAGAAAAAGAAAGGCGTCT
19AH CCTTGAAACCATCTGAAATTCCTGATCCTTTTACAGAGCAAGAGAAAAAGAAAGGCGTCT
23FPO CCTTGAAACCATCTGAAATTCCTGATCCTTTTACAGAGCAAGAGAAAAAGAAAGGCGTCT
19FTW CCTTGAAACCATCTGAAATTCCTGATCCTTTTACAGATCAGGAAAAGAACAGGGAGTTT
9VSP CCTTGAAACCATCTGAAATTCCTGATCCTTTTACAGAGCAAGAGAAAAAGAAAGGCGTCT
TIGR4 CCTTGAAACCATCTGAAATTCCTGATCCTTTTACAGAGCAAGAGAAAAAGAAAGGCGTCT
23FTW CCTTGAAACCATCTGAAATTCCTGATCCTTTTACAGAGCAAGAGAAAAAGAAAGGCGTCT

14CSR CAGAATATGCCAATATGCTAAAGGTCCATGAGCGGATTGGCTATGTGGAAATTCCTGCGA
670 CAGAATATGCCAATATGCTAAAGGTCCATGAGCGGATTGGCTATGTGGAAATTCCTGCGA
6BF CAGAATATGCCAATATGCTAAAGGTCCATGAGCGGATTGGCTATGTGGAAATTCCTGCGA
6BSP CAGAATATGCCAATATGCTAAAGGTCCATGAGCGGATTGGCTATGTGGAAATTCCTGCGA
19AH CAGAATATGCCAATATGCTAAAGGTCCATGAGCGGATTGGCTATGTGGAAATTCCTGCGA
23FPO CAGAATATGCCAATATGCTAAAGGTCCATGAGCGGATTGGCTATGTGGAAATTCCTGCGA
19FTW CAGAATATGCTAACATGCTAAAGTTCATGAGCGTATCGGATATGTAGAAATTCCTGCGA
9VSP CAGAATATGCCAATATGCTAAAGGTCCATGAGCGGATTGGCTATGTGGAAATTCCTGCGA
TIGR4 CAGAATATGCCAATATGCTAAAGGTCCATGAGCGGATTGGCTATGTGGAAATTCCTGCGA
23FTW CAGAATATGCCAATATGCTAAAGGTCCATGAGCGGATTGGCTATGTGGAAATTCCTGCGA

14CSR TTGATCAGGAAATTCGGATGTATGTGCGAACGAGTGAGGAAATTCCTCAGAAGGGCGCAG
670 TTGATCAGGAAATTCGGATGTATGTGCGAACGAGTGAGGAAATTCCTCAGAAGGGCGCAG
6BF TTGATCAGGAAATTCGGATGTATGTGCGAACGAGTGAGGAAATTCCTCAGAAGGGCGCAG
6BSP TTGATCAGGAAATTCGGATGTATGTGCGAACGAGTGAGGAAATTCCTCAGAAGGGCGCAG
19AH TTGATCAGGAAATTCGGATGTATGTGCGAACGAGTGAGGAAATTCCTCAGAAGGGCGCAG
23FPO TTGATCAGGAAATTCGGATGTATGTGCGAACGAGTGAGGAAATTCCTCAGAAGGGCGCAG
19FTW TTGAACAGGAAATCCCATGTATGTTGGCACAGTGAAGACATTCCTCAGAAAGGGCGCAG
9VSP TTGATCAGGAAATTCGGATGTATGTGCGAACGAGTGAGGAAATTCCTCAGAAGGGCGCAG
TIGR4 TTGATCAGGAAATTCGGATGTATGTGCGAACGAGTGAGGAAATTCCTCAGAAGGGCGCAG
23FTW TTGATCAGGAAATTCGGATGTATGTGCGAACGAGTGAGGAAATTCCTCAGAAGGGCGCAG

14CSR GATTGCTAGAGGGAGCTTCGTTACCGGTTGGTGGTGAAAATACCCACACAGTTGTCACTG
670 GATTGCTAGAGGGAGCTTCGTTACCGGTTGGTGGTGAAAATACCCACACAGTTGTCACTG
6BF GATTGCTAGAGGGAGCTTCGTTACCGGTTGGTGGTGAAAATACCCACACAGTTGTCACTG
6BSP GATTGCTAGAGGGAGCTTCGTTACCGGTTGGTGGTGAAAATACCCACACAGTTGTCACTG
19AH GATTGCTAGAGGGAGCTTCGTTACCGGTTGGTGGTGAAAATACCCACACAGTTGTCACTG
23FPO GATTGCTAGAGGGAGCTTCGTTACCGGTTGGTGGTGAAAATACCCACACAGTTGTCACTG
19FTW GGCTGTTAGAAGGGGCTTCGCTGCCTGTTGGAGGTGAAAATACCCATACAGTGATCACTG
9VSP GATTGCTAGAGGGAGCTTCGTTACCGGTTGGTGGTGAAAATACCCACACAGTTGTCACTG
TIGR4 GGCTGTTAGAAGGGGCTTCGCTGCCTGTTGGAGGTGAAAATACCCATACAGTGATCACTG
23FTW GATTGCTAGAGGGAGCTTCGTTACCGGTTGGTGGTGAAAATACCCACACAGTTGTCACTG
* * * * *

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14CSR CTCATAGAGGATTACCGACGGCAGAACTGTTTAGTCAATTGGATAAGATGAAAAAAGGGG
670 CTCATAGAGGATTACCGACGGCAGAACTGTTTAGTCAATTGGATAAGATGAAAAAAGGGG
6BF CTCATAGAGGATTACCGACGGCAGAACTGTTTAGTCAATTGGATAAGATGAAAAAAGGGG
6BSP CTCATAGAGGATTACCGACGGCAGAACTGTTTAGTCAATTGGATAAGATGAAAAAAGGGG
19AH CTCATAGAGGATTACCGACGGCAGAACTGTTTAGTCAATTGGATAAGATGAAAAAAGGGG
23FPO CTCATAGAGGATTACCGACGGCAGAACTGTTTAGTCAATTGGATAAGATGAAAAAAGGGG
19FTW CTCACAGAGGATTGCCAACGGCAGAACTGTTTCACTCAATTGGATAAGATGAAGAAAGGGG
9VSP CTCACAGAGGATTGCCAACGGCAGAACTGTTTAGTCAATTGGATAAGATGAAAAAAGGGG
TIGR4 CTCACAGAGGATTGCCAACGGCAGAACTGTTTCACTCAATTGGATAAGATGAAAAAAGGGG
23FTW CTCATAGAGGATTACCGACGGCAGAACTGTTTAGTCAATTGGATAAGATGAAAAAAGGGG

14CSR ATGTCTTTTATCTTCACGTTTTAGACCAGGTGTTGGCCTACCAAGTGGATCAGATTTTGA
670 ATGTCTTTTATCTTCACGTTTTAGACCAGGTGTTGGCCTACCAAGTGGATCAGATTTTGA
6BF ATGTCTTTTATCTTCACGTTTTAGACCAGGTGTTGGCCTACCAAGTGGATCAGATTTTGA
6BSP ATGTCTTTTATCTTCACGTTTTAGACCAGGTGTTGGCCTACCAAGTGGATCAGATTTTGA
19AH ATGTCTTTTATCTTCACGTTTTAGACCAGGTGTTGGCCTACCAAGTGGATCAGATTTTGA
23FPO ATATCTTTTATCTTCACGTTTTAGATCAGGTGTTGGCCTACCAAGTGGATCAGATAGTGA
19FTW ATATCTTTTATCTTCACGTTTTAGACCAGGTGTTGGCCTATCAAGTGGATCAGATAGTGA
9VSP ATATCTTTTATCTTCACGTTTTAGATCAGGTGTTGGCCTACCAAGTGGATCAGATAGTGA
TIGR4 ATATCTTTTATCTTCACGTTTTAGATCAGGTGTTGGCCTACCAAGTGGATCAGATAGTGA
23FTW ATGTCTTTTATCTTCACGTTTTAGACCAGGTGTTGGCCTACCAAGTGGATCAGATTTTGA
** *****

14CSR CGGTTGAGCCAAATGACTTTGAGCCTGTCTTGATTCAACATGGGGAAGATTATGCGACCT
670 CGGTTGAGCCAAATGACTTTGAGCCTGTCTTGATTCAACATGGGGAAGATTATGCGACCT
6BF CGGTTGAGCCAAATGACTTTGAGCCTGTCTTGATTCAACATGGGGAAGATTATGCGACCT
6BSP CGGTTGAGCCAAATGACTTTGAGCCTGTCTTGATTCAACATGGGGAAGATTATGCGACCT
19AH CGGTTGAGCCAAATGACTTTGAGCCTGTCTTGATTCAACATGGGGAAGATTATGCGACCT
23FPO CGGTTGAGCCGAATGACTTTGAGCCTGTCTTGATTCAACATGGGGAAGATTATGCGACCT
19FTW CGGTTGAGCCGAATGATTTTGAAGCCTGTCTTGATTCAACATGGGGAAGATTATGCGACCT
9VSP CGGTTGAGCCGAATGACTTTGAGCCTGTCTTGATTCAACATGGGGAAGATTATGCGACCT
TIGR4 CGGTTGAGCCGAATGACTTTGAGCCTGTCTTGATTCAACATGGGGAAGATTATGCGACCT
23FTW CGGTTGAGCCAAATGACTTTGAGCCTGTCTTGATTCAACATGGGGAAGATTATGCGACCT

14CSR TGTTGACCTGTACACCGTATATGATTAACAGTCATCGTCTGTTGGTACGTGGGAAGCGGA
670 TGTTGACCTGTACACCGTATATGATTAACAGTCATCGTCTGTTGGTACGTGGGAAGCGGA
6BF TGTTGACCTGTACACCGTATATGATTAACAGTCATCGTCTGTTGGTACGTGGGAAGCGGA
6BSP TGTTGACCTGTACACCGTATATGATTAACAGTCATCGTCTGTTGGTACGTGGGAAGCGGA
19AH TGTTGACCTGTACACCGTATATGATTAACAGTCATCGTCTGTTGGTACGTGGGAAGCGGA
23FPO TGTTGACTTGTAACCGTATATGATTAACAGTCATCGTCTGTTGGTACGTGGGAAGCGGA
19FTW TACTGACTTGTAACCGTATATGATTAACAGTCATCGTCTGTTGGTACGTGGGAAGCGGA
9VSP TGTTGACTTGTAACCGTATATGATTAACAGTCATCGTCTGTTGGTACGTGGGAAGCGGA
TIGR4 TGTTGACTTGTAACCGTATATGATTAACAGTCATCGTCTGTTGGTACGTGGGAAGCGGA
23FTW TGTTGACCTGTACACCGTATATGATTAACAGTCATCGTCTGTTGGTACGTGGGAAGCGGA
* *****

14CSR TTCCGTATACGGCACCAATTGCAGAGCGAAATCGAGCGGTGAGAGAGCGTGGGCAATTCT
670 TTCCGTATACGGCACCAATTGCAGAGCGAAATCGAGCGGTGAGAGAGCGTGGGCAATTCT
6BF TTCCGTATACGGCACCAATTGCAGAGCGAAATCGAGCGGTGAGAGAGCGTGGGCAATTCT
6BSP TTCCGTATACGGCACCAATTGCAGAGCGAAATCGAGCGGTGAGAGAGCGTGGGCAATTCT
19AH TTCCGTATACGGCACCAATTGCAGAGCGAAATCGAGCGGTGAGAGAGCGTGGGCAATTCT
23FPO TTCCGTATACGGCACCAATTGCAGAGCGAAATCGAGCGGTGAGAGAGCGTGGGCAATTCT
19FTW TTCCATATACAGCGCGGATTGCTGAGCGGAATCGAGCGGTGAGAGAGCGTGGGCAATTCT
9VSP TTCCGTATACGGCACCAATTGCAGAGCGAAATCGAGCGGTGAGAGAGCGTGGGCAATTCT
TIGR4 TTCCGTATACGGCACCAATTGCAGAGCGAAATCGAGCGGTGAGAGAGCGTGGGCAATTCT
23FTW TTCCGTATACGGCACCAATTGCAGAGCGAAATCGAGCGGTGAGAGAGCGTGGGCAATTCT

Figure 196I

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14CSR GGTGTGGTTATTGCTAGCGGCGTTGGTTATGATTCTGGTATTGAGTTACGGGGTGTATC
670 GGTGTGGTTATTGCTAGCGGCGTTGGTTATGATTCTGGTATTGAGTTACGGGGTGTATC
6BF GGTGTGGTTATTGCTAGCGGCGTTGGTTATGATTCTGGTATTGAGTTACGGGGTGTATC
6BSP GGTGTGGTTATTGCTAGCGGCGTTGGTTATGATTCTGGTATTGAGTTACGGGGTGTATC
19AH GGTGTGGTTATTGCTAGCGGCGTTGGTTATGATTCTGGTATTGAGTTACGGGGTGTATC
23FPO GGTGTGGTTATTACTAGGAGCGATGGCGGTATCCTTCTCTTGTGTATCGCGTGTATC
19FTW GGTGTGGTTATTACTAGGAGCGATGGCGGTATCCTTCTCTTGTGTATCGCGTGTATC
9VSP GGTGTGGTTATTACTAGGAGCGATGGCGGTATCCTTCTCTTGTGTATCGCGTGTATC
TIGR4 GGTGTGGTTATTACTAGGAGCGATGGCGGTATCCTTCTCTTGTGTATCGCGTGTATC
23FTW GGTGTGGTTATTGCTAGCGGCGTTGGTTATGATTCTGGTATTGAGTTACGGGGTGTATC

14CSR GTCATCGTCGCATTGTCAAAGGGCTAGAAAAACAATTGGAGGAGCATCATGTCAAAGGCT
670 GTCATCGTCGCATTGTCAAAGGGCTAGAAAAACAATTGGAGGAGCATCATGTCAAAGGCT
6BF GTCATCGTCGCATTGTCAAAGGGCTAGAAAAACAATTGGAGGAGCATCATGTCAAAGGCT
6BSP GTCATCGTCGCATTGTCAAAGGGCTAGAAAAACAATTGGAGGAGCATCATGTCAAAGGCT
19AH GTCATCGTCGCATTGTCAAAGGGCTAGAAAAACAATTGGAGGAGCATCATGTCAAAGGCT
23FPO GTAATCGACGGATTGTCAAAGGACTAGAAAAGCAATTGGAGGGGCGTCATGTCAAAGGACT
19FTW GTAATCGACGGATTGTCAAAGGACTAGAAAAGCAATTGGAGGGGCGTCATGTCAAAGGACT
9VSP GTAATCGACGGATTGTCAAAGGACTAGAAAAGCAATTGGAGGGGCGTCATGTCAAAGGACT
TIGR4 GTAATCGACGGATTGTCAAAGGACTAGAAAAGCAATTGGAGGGGCGTCATGTCAAAGGACT
23FTW GTCATCGTCGCATTGTCAAAGGGCTAGAAAAACAATTGGAGGAGCATCATGTCAAAGGCT
** * * * *

14CSR AAGCTACAGAAATTACTAGGGTATTTGCTGATGCTGGTAGCATTGGTGATTCCTGTTTAT
670 AAGCTACAGAAATTACTAGGGTATTTGCTGATGCTGGTAGCATTGGTGATTCCTGTTTAT
6BF AAGCTACAGAAATTACTAGGGTATTTGCTGATGCTGGTAGCATTGGTGATTCCTGTTTAT
6BSP AAGCTACAGAAATTACTAGGGTATTTGCTGATGCTGGTAGCATTGGTGATTCCTGTTTAT
19AH AAGCTACAGAAATTACTAGGGTATTTGCTGATGCTGGTAGCATTGGTGATTCCTGTTTAT
23FPO AAATACGAGCCTTATTGGGATACTTGTGATGTTGGTAGCCTGTTTGATTCCTATTTAT
19FTW AAATACGAGCCTTATTGGGATACTTGTGATGTTGGTAGCCTGTTTGATTCCTATTTAT
9VSP AAATACGAGCCTTATTGGGATACTTGTGATGTTGGTAGCCTGTTTGATTCCTATTTAT
TIGR4 AAATACGAGCCTTATTGGGATACTTGTGATGTTGGTAGCCTGTTTGATTCCTATTTAT
23FTW AAGCTACAGAAATTACTAGGGTATTTGCTGATGCTGGTAGCATTGGTGATTCCTGTTTAT
** * * *

14CSR TGTTTTGGGCAGATGGTGTTACAGTCTTTAGGACAAGTAAAAGGTCATGAGATATTTTCA
670 TGTTTTGGGCAGATGGTGTTACAGTCTTTAGGACAAGTAAAAGGTCATGAGATATTTTCA
6BF TGTTTTGGGCAGATGGTGTTACAGTCTTTAGGACAAGTAAAAGGTCATGAGATATTTTCA
6BSP TGTTTTGGGCAGATGGTGTTACAGTCTTTAGGACAAGTAAAAGGTCATGAGATATTTTCA
19AH TGTTTTGGGCAGATGGTGTTACAGTCTTTAGGACAAGTAAAAGGTCATGAGATATTTTCA
23FPO TGTTTTGGGCAGATGGTGTTGACAGTCTCTTGGACAGGTGAAAGGTCATGCTACATTTGTG
19FTW TGTTTTGGGCAGATGGTGTTGACAGTCTCTTGGACAGGTGAAAGGTCATGCTACATTTGTG
9VSP TGTTTTGGGCAGATGGTGTTGACAGTCTCTTGGACAGGTGAAAGGTCATGCTACATTTGTG
TIGR4 TGTTTTGGGCAGATGGTGTTGACAGTCTCTTGGACAGGTGAAAGGTCATGCTACATTTGTG
23FTW TGTTTTGGGCAGATGGTGTTACAGTCTTTAGGACAAGTAAAAGGTCATGAGATATTTTCA

14CSR GAATCTGTGACGGCCGACAGTTACCAAGAGCAATTGCAACGGTCGCTTGATTACAATCAA
670 GAATCTGTGACGGCCGACAGTTACCAAGAGCAATTGCAACGGTCGCTTGATTACAATCAA
6BF GAATCTGTGACGGCCGACAGTTACCAAGAGCAATTGCAACGGTCGCTTGATTACAATCAA
6BSP GAATCTGTGACGGCCGACAGTTACCAAGAGCAATTGCAACGGTCGCTTGATTACAATCAA
19AH GAATCTGTGACGGCCGACAGTTACCAAGAGCAATTGCAACGGTCGCTTGATTACAATCAA
23FPO AAATCCATGACAACCTGAAATGTACCAAGAACAACAGAACCATTCTCTCGCCTACAATCAA
19FTW AAATCCATGACAACCTGAAATGTACCAAGAACAACAGAACCATTCTCTCGCCTACAATCAA
9VSP AAATCCATGACAACCTGAAATGTACCAAGAACAACAGAACCATTCTCTCGCCTACAATCAA
TIGR4 AAATCCATGACAACCTGAAATGTACCAAGAACAACAGAACCATTCTCTCGCCTACAATCAA
23FTW GAATCTGTGACGGCCGACAGTTACCAAGAGCAATTGCAACGGTCGCTTGATTACAATCAA

Figure 196AJ

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14CSR      CGCTTGGATTTCGCAAAATCGTATTGTAGATCCTTTTTTGGCGGAAGGGTATGAGGTAAAT
670        CGCTTGGATTTCGCAAAATCGTATTGTAGATCCTTTTTTGGCGGAAGGGTATGAGGTAAAT
6BF        CGCTTGGATTTCGCAAAATCGTATTGTAGATCCTTTTTTGGCGGAAGGGTATGAGGTAAAT
6BSP       CGCTTGGATTTCGCAAAATCGTATTGTAGATCCTTTTTTGGCGGAAGGGTATGAGGTAAAT
19AH       CGCTTGGATTTCGCAAAATCGTATTGTAGATCCTTTTTTGGCGGAAGGGTATGAGGTAAAT
23FPO      CGCTTGGCTTCGCAAAATCGCATTGTAGATCCTTTTTTGGCGGAGGGATATGAGGTCAAT
19FTW      CGCTTGGCTTCGCAAAATCGCATTGTAGATCCTTTTTTGGCGGAGGGATATGAGGTCAAT
9VSP       CGCTTGGCTTCGCAAAATCGCATTGTAGATCCTTTTTTGGCGGAGGGATATGAGGTCAAT
TIGR4      CGCTTGGCTTCGCAAAATCGCATTGTAGATCCTTTTTTGGCGGAGGGATATGAGGTCAAT
23FTW      CGCTTGGATTTCGCAAAATCGTATTGTAGATCCTTTTTTGGCGGAAGGGTATGAGGTAAAT
*****

14CSR      TACCAAGTGTCTGACGATCCTGATGCAGTCTACGGCTATTTGTCGATTCCGAGTTTGGAA
670        TACCAAGTGTCTGACGATCCTGATGCAGTCTACGGCTATTTGTCGATTCCGAGTTTGGAA
6BF        TACCAAGTGTCTGACGATCCTGATGCAGTCTACGGCTATTTGTCGATTCCGAGTTTGGAA
6BSP       TACCAAGTGTCTGACGATCCTGATGCAGTCTACGGCTATTTGTCGATTCCGAGTTTGGAA
19AH       TACCAAGTGTCTGACGATCCTGATGCAGTCTACGGCTATTTGTCGATTCCGAGTTTGGAA
23FPO      TACCAAGTGTCTGACGACCTGATGCAGTCTATGGTTACTTGTCTATTCCAAGTTTGGAA
19FTW      TACCAAGTGTCTGACGACCTGATGCAGTCTATGGTTACTTGTCTATTCCAAGTTTGGAA
9VSP       TACCAAGTGTCTGACGACCTGATGCAGTCTATGGTTACTTGTCTATTCCAAGTTTGGAA
TIGR4      TACCAAGTGTCTGACGACCTGATGCAGTCTATGGTTACTTGTCTATTCCAAGTTTGGAA
23FTW      TACCAAGTGTCTGACGATCCTGATGCAGTCTACGGCTATTTGTCGATTCCGAGTTTGGAA
*****

14CSR      ATCATGGAGCCAGTTTATCTAGGAGCGGATTACCATCATTTAGCAATGGGGTTGGCCCAT
670        ATCATGGAGCCAGTTTATCTAGGAGCGGATTACCATCATTTAGCAATGGGGTTGGCCCAT
6BF        ATCATGGAGCCAGTTTATCTAGGAGCGGATTACCATCATTTAGCAATGGGGTTGGCCCAT
6BSP       ATCATGGAGCCAGTTTATCTAGGAGCGGATTACCATCATTTAGCAATGGGGTTGGCCCAT
19AH       ATCATGGAGCCAGTTTATCTAGGAGCGGATTACCATCATTTAGCAATGGGGTTGGCCCAT
23FPO      ATCATGGAGCCGTTTATTTGGGAGCAGATTATCATCATTTAGGGATGGGCTTGGCTCAT
19FTW      ATCATGGAGCCGTTTATTTGGGAGCAGATTATCATCATTTAGGGATGGGCTTGGCTCAT
9VSP       ATCATGGAGCCGTTTATTTGGGAGCAGATTATCATCATTTAGGGATGGGCTTGGCTCAT
TIGR4      ATCATGGAGCCGTTTATTTGGGAGCAGATTATCATCATTTAGGGATGGGCTTGGCTCAT
23FTW      ATCATGGAGCCAGTTTATCTAGGAGCGGATTACCATCATTTAGCAATGGGGTTGGCCCAT
*****

14CSR      GTGGATGGGACGCCTCTTCTGTTGAGGGAAAAGGGATTTCGTTCACTGATTGCTGGGCAC
670        GTGGATGGGACGCCTCTTCTGTTGAGGGAAAAGGGATTTCGTTCACTGATTGCTGGGCAC
6BF        GTGGATGGGACGCCTCTTCTGTTGAGGGAAAAGGGATTTCGTTCACTGATTGCTGGGCAC
6BSP       GTGGATGGGACGCCTCTTCTGTTGAGGGAAAAGGGATTTCGTTCACTGATTGCTGGGCAC
19AH       GTGGATGGGACGCCTCTTCTGTTGAGGGAAAAGGGATTTCGTTCACTGATTGCTGGGCAC
23FPO      GTGGATGGTACACCGCTGCCTCTGGATGGTACAGGGATTTCGCTCAGTGATTGCTGGGCAC
19FTW      GTGGATGGTACACCGCTGCCTCTGGATGGTACAGGGATTTCGCTCAGTGATTGCTGGGCAC
9VSP       GTGGATGGTACACCGCTGCCTCTGGATGGTACAGGGATTTCGCTCAGTGATTGCTGGGCAC
TIGR4      GTGGATGGTACACCGCTGCCTCTGGATGGTACAGGGATTTCGCTCAGTGATTGCTGGGCAC
23FTW      GTGGATGGGACGCCTCTTCTGTTGAGGGAAAAGGGATTTCGTTCACTGATTGCTGGGCAC
*****

14CSR      CGTGCAGAACCAAGCCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCT
670        CGTGCAGAACCAAGCCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCT
6BF        CGTGCAGAACCAAGCCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCT
6BSP       CGTGCAGAACCAAGCCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCT
19AH       CGTGCAGAACCAAGCCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCT
23FPO      CGTGCAGAGCCAAGCCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCT
19FTW      CGTGCAGAGCCAAGCCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCT
9VSP       CGTGCAGAGCCAAGCCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCT
TIGR4      CGTGCAGAGCCAAGCCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCT
23FTW      CGTGCAGAACCAAGCCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCT
*****
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Figure 196AK

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14CSR CTTTATTATGATAATGGCCAGGAAATTGTAGAATATCAGATGATGGACACAGAGATTATT
670 CTTTATTATGATAATGGCCAGGAAATTGTAGAATATCAGATGATGGACACAGAGATTATT
6BF CTTTATTATGATAATGGCCAGGAAATTGTAGAATATCAGATGATGGACACAGAGATTATT
6BSP CTTTATTATGATAATGGCCAGGAAATTGTAGAATATCAGATGATGGACACAGAGATTATT
19AH CTTTATTATGATAATGGCCAGGAAATTGTAGAATATCAGATGATGGACACAGAGATTATT
23FPO CTTTATTATGATAATGGCCAGGAAATTGTAGAATATCAGATGATGGACACAGAGATTATT
19FTW CTTTATTATGATAATGGCCAGGAAATTGTAGAATATCAGATGATGGACACAGAGATTATT
9VSP CTTTATTATGATAATGGCCAGGAAATTGTAGAATATCAGATGATGGACACAGAGATTATT
TIGR4 CTTTATTATGATAATGGCCAGGAAATTGTAGAATATCAGATGATGGACACAGAGATTATT
23FTW CTTTATTATGATAATGGCCAGGAAATTGTAGAATATCAGATGATGGACACAGAGATTATT

14CSR TTACCGTCGGAATGGGAAAAATTAGAATCGGTTAGCTCTAAAAATATCATGACCTTGATA
670 TTACCGTCGGAATGGGAAAAATTAGAATCGGTTAGCTCTAAAAATATCATGACCTTGATA
6BF TTACCGTCGGAATGGGAAAAATTAGAATCGGTTAGCTCTAAAAATATCATGACCTTGATA
6BSP TTACCGTCGGAATGGGAAAAATTAGAATCGGTTAGCTCTAAAAATATCATGACCTTGATA
19AH TTACCGTCGGAATGGGAAAAATTAGAATCGGTTAGCTCTAAAAATATCATGACCTTGATA
23FPO TTACCGTCGGAATGGGAAAAATTAGAATCGGTTAGCTCTAAAAATATCATGACCTTGATA
19FTW TTACCGTCGGAATGGGAAAAATTAGAATCGGTTAGCTCTAAAAATATCATGACCTTGATA
9VSP TTACCGTCGGAATGGGAAAAATTAGAATCGGTTAGCTCTAAAAATATCATGACCTTGATA
TIGR4 TTACCGTCGGAATGGGAAAAATTAGAATCGGTTAGCTCTAAAAATATCATGACCTTGATA
23FTW TTACCGTCGGAATGGGAAAAATTAGAATCGGTTAGCTCTAAAAATATCATGACCTTGATA

14CSR ACCTGCGATCCGATTCCCTACCTTTAATAAACGCTTATTAGTGAATTTTGAACGAGTCGCT
670 ACCTGCGATCCGATTCCCTACCTTTAATAAACGCTTATTAGTGAATTTTGAACGAGTCGCT
6BF ACCTGCGATCCGATTCCCTACCTTTAATAAACGCTTATTAGTGAATTTTGAACGAGTCGCT
6BSP ACCTGCGATCCGATTCCCTACCTTTAATAAACGCTTATTAGTGAATTTTGAACGAGTCGCT
19AH ACCTGCGATCCGATTCCCTACCTTTAATAAACGCTTATTAGTGAATTTTGAACGAGTCGCT
23FPO ACCTGCGATCCGATTCCCTACCTTTAATAAACGCTTATTAGTGAATTTTGAACGAGTCGCT
19FTW ACCTGCGATCCGATTCCCTACCTTTAATAAACGCTTATTAGTGAATTTTGAACGAGTCGCT
9VSP ACCTGCGATCCGATTCCCTACCTTTAATAAACGCTTATTAGTGAATTTTGAACGAGTCGCT
TIGR4 ACCTGCGATCCGATTCCCTACCTTTAATAAACGCTTATTAGTGAATTTTGAACGAGTCGCT
23FTW ACCTGCGATCCGATTCCCTACCTTTAATAAACGCTTATTAGTGAATTTTGAACGAGTCGCT

14CSR GTTTATCAAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTTTACGAAAGAA
670 GTTTATCAAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTTTACGAAAGAA
6BF GTTTATCAAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTTTACGAAAGAA
6BSP GTTTATCAAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTTTACGAAAGAA
19AH GTTTATCAAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTTTACGAAAGAA
23FPO GTTTATCAAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTTTACGAAAGAA
19FTW GTTTATCAAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTTTACGAAAGAA
9VSP GTTTATCAAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTTTACGAAAGAA
TIGR4 GTTTATCAAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTTTACGAAAGAA
23FTW GTTTATCAAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTTTACGAAAGAA

14CSR GGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTGTACCGTGGGCTAGTGGTACTG
670 GGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTGTACCGTGGGCTAGTGGTACTG
6BF GGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTGTACCGTGGGCTAGTGGTACTG
6BSP GGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTGTACCGTGGGCTAGTGGTACTG
19AH GGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTGTACCGTGGGCTAGTGGTACTG
23FPO GGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTGTACCGTGGGCTAGTGGTACTG
19FTW GGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTGTACCGTGGGCTAGTGGTACTG
9VSP GGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTGTACCGTGGGCTAGTGGTACTG
TIGR4 GGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTGTACCGTGGGCTAGTGGTACTG
23FTW GGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTGTACCGTGGGCTAGTGGTACTG

Figure 196AL

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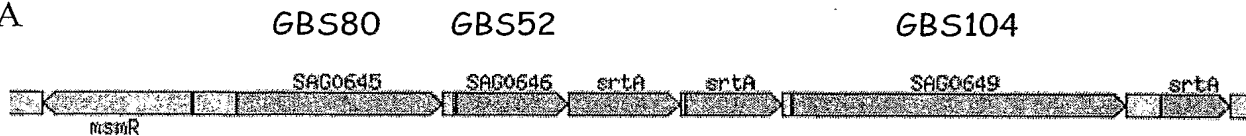
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14CSR      GCATTTCTGGGAATCCTGTTTGTGTTTGTGGAAGCTAGCACGTTTACTACGAGGGAAATAA
670        GCATTTCTGGGAATCCTGTTTGTGTTTGTGGAAGCTAGCACGTTTACTACGAGGGAAATAA
6BF        GCATTTCTGGGAATCCTGTTTGTGTTTGTGGAAGCTAGCACGTTTACTACGAGGGAAATAA
6BSP       GCATTTCTGGGAATCCTGTTTGTGTTTGTGGAAGCTAGCACGTTTACTACGAGGGAAATAA
19AH       GCATTTATGGGAATCCTGTTTGTGTTTGTGGAAGCTAGCACGTTTACTACGAGGGAAATAA
23FPO      GCATTTCTGGGAATCCTGTTTGTGTTTGTGGAAGCTAGCACGTTTACTACGAGGGAAATAA
19FTW      GCATTTCTGGGAATCCTGTTTGTGTTTGTGGAAGCTAGCACGTTTACTACGAGGGAAATAA
9VSP       GCATTTCTGGGAATCCTGTTTGTGTTTGTGGAAGCTAGCACGTTTACTACGAGGGAAATAA
TIGR4      GCATTTCTGGGAATCCTGTTTGTGTTTGTGGAAGCTAGCACGTTTACTACGAGGGAAATAA
23FTW      GCATTTCTGGGAATCCTGTTTGTGTTTGTGGAAGCTAGCACGTTTACTACGAGGGAAATAA
          *****

14CSR      AAAGAAATGAAAGGAAAGCTAAGGCTGTTCCCTTTTTCCGGCTCTTTGTCAACTGTAGGGG
670        AAAGAAATGAAAGGAAAGCTAAGGCTGTTCCCTTTTTCCGGCTCTTTGTCAACTGTAGTGG
6BF        AAAGAAATGAAAGGAAAGCTAAGGCTGTTCCCTTTTTCCGGCTCTTTGTCAACTGTAG---
6BSP       AAAGAAATGAAAGGAAAGCTAAGGCTGTTCCCTTTTTCCGGCTCTTTGTCAACTGTAG---
19AH       AAAGAAATGAAAGGAAAGCTAAGGCTGTTCCCTTTTTCCGGCTCTTTGTCAACTGTAG---
23FPO      AAAGAAATGAAAGGAAAGCTAAGGCTGTTCCCTTTTTCCGGCTCTTTGTCAACTGT-----
19FTW      AAAGAAATGAAAGGAAAGCTAAGGCTGTTCCCTTTTTCCGGCTCTTTGTCAACTGTAGT--
9VSP       AAAGAAATGAAAGGAAAGCTAAGGCTGTTCCCTTTTTCCGGCTCTTTGTCAACTGTAG---
TIGR4      AAAGAAATGAAAGGAAAGCTAAGGCTGTTCCCTTTTTCCGGCTCTTTGTCAACTGTAGTGG
23FTW      AAAGAAATGAAAGGAAAGCTAAGGCTGTTCCCTTTTTCCGGCTCTTTGTCAACTGTAG---
          *****
```

Figure 196AM

Figure 197

A



B

Intergenic region between AraC R and GBS 80

AraC...CAT

TTGATAGAC**CCGCCTTC**ATTATCATTTCTAGAATTTTTCTTTAGGTTTGTA
 AAGACTACAAAATAAAATGATGAAAACAACATCTTGTGGATACACTAAA
 AAGACACGCTAATTAGCAAACCTCTCTTCATCATCTCTCACCATTATTA
 TACTAC **TATTTATAT**GACAAATAAAGGT**GATTT** **TGTTAA**AATATAACTTT
 GAAAATCCACATATATTTTTTAATCTTCCGTCTG**AAAAAA**TAAATAAAAAT
 AGTAAAAATAAACACGAATTTAAAATAAGCAAATTTTTTAAGAAAATCTG
 TGCTAAACTTTAATAGTTTTGTGCTTAATAATAATCAGCACTTACAAAGA
 ACAAAGGGAAAAGCGAG**GAGAG**AACTTTTA **ATG... GBS80**

C

187	4A		5A		5A
233	6A		6A		7A
Strain	FACS α -80	Strain	FACS α -80	Strain	FACS α -80
1998	95	5364	454	2129	57
2110	0	JMV071	556	2274	113
2603	62	JM91003	587	5401	170
3050	43	CJB111	365	5408	0
5376	165			5518	31
M781	65			CJB110	71
COH1	305 (G→T 179)			J7357B	91
18rs 21	0 (STOP, no LPXTG)			COH31	0

PCT/US05/2723433/487

AI-1											
			aa	M1	M3	M5	M18	M49	M6	M12	
M6											
50913503	M6_Spy0157	LPXTG	628	gas15 30%in593aa	M3-0098 46%in256aa M3-0104 28%in563aa		M18-0132 24%in701aa			M12-4134 74%in703aa	Fibronecti n-binding protein (protein F)
50913505	M6_Spy0159	LPXSG	1037		M3-0104 25%in339aa					M12-4141 37%in98aa	Collagen adhesion protein
50913506	M6_Spy0160	LPXTG	557								Fimbrial structural subunit

Figure 198

AI-2									
		aa	M1	M3	M5	M18	M49	M6	M12
M1									
gas15	gas15	VXGTG	762	M3-0098 50%in738aa	M5-orf78 60%in462aa	M18-0126 54%in469aa			M12-4135 54%in747aa
13621428	SPy0128 gas16	EVXTG	340	M3-0100 40%in354aa	M5-orf80 41%in358aa	M18-0128 38%in357aa			M12-4137 40%in354aa
13621430	SPy0130 gas18	LPXTG	215	M3-0102 32%200aa	M5-orf82 31%in213aa	M18-0130 32%in213aa			M12-4139 31%in206aa
									hypothetical protein (fimbrial)
									hypothetical protein

Figure 199

AI-3										
			aa	M1	M3	M5	M18	M49	M6	M12
M3										
21909634	SpyM3_0098	VPXTG	744	gas15 51%in739aa		M5-orf78 58%in484aa	M18-0126 74%in482aa			M12-4135 55%in751aa
21909636	SpyM3_0100	QVXTG	344	gas16 40%in354aa		M5-orf80 64%in349aa	M18-0128 67%in345aa			M12-4137 61%in344aa
21909638	SpyM3_0102	LPXAG	195	gas18 32%in200aa		M5-orf82 98%in183aa	M18-0130 97%in183aa			M12-4139 99%in183aa
21909640	SpyM3_0104	LPXTG	696			M5-orf84 88%in656aa	M18-0132 88%in656aa			M12-4141 59%in612aa
										putative collagen binding protein (Cpb)
										conserved hypothetic al protein (fimbrial)
										hypothetic al protein
										protein F2 like fibronectin -binding

Figure 200A

Figure 200B

Figure 200B

Figure 200C

Figure 200C

M49												
56808848	VPXTG	744	gas15 55%in738aa	M3-0098 72%in743aa	M5-orf78 78%in483	M18-0126 61% in484				M12-4135 73%in752aa	putative collagen binding protein (Cpb)	
56808846	QVXTG	344	gas16 36%in355aa	M3-0100 66%in345aa	M5-orf80 61%in349aa	M18-0128 90%in344aa				M12-4137 62%in344aa	conserved hypothetic al protein (fimbrial)	
56808844	LPXAG	189	gas18 31%in206aa	M3-102 98%in189aa	M5-orf82 98%in189aa	M18-0130 98%in189aa				M12-4139 98%in189aa	hypothetic al protein	
56808842	LPXTG	1160		M3-104 59%in612aa	M5-orf84 50%in701aa	M18-0132 50%in701aa			M6-0157 32%in296aa	M12-4141 91%in1164aa	protein F2 like fibronectin -binding	

Figure 200D

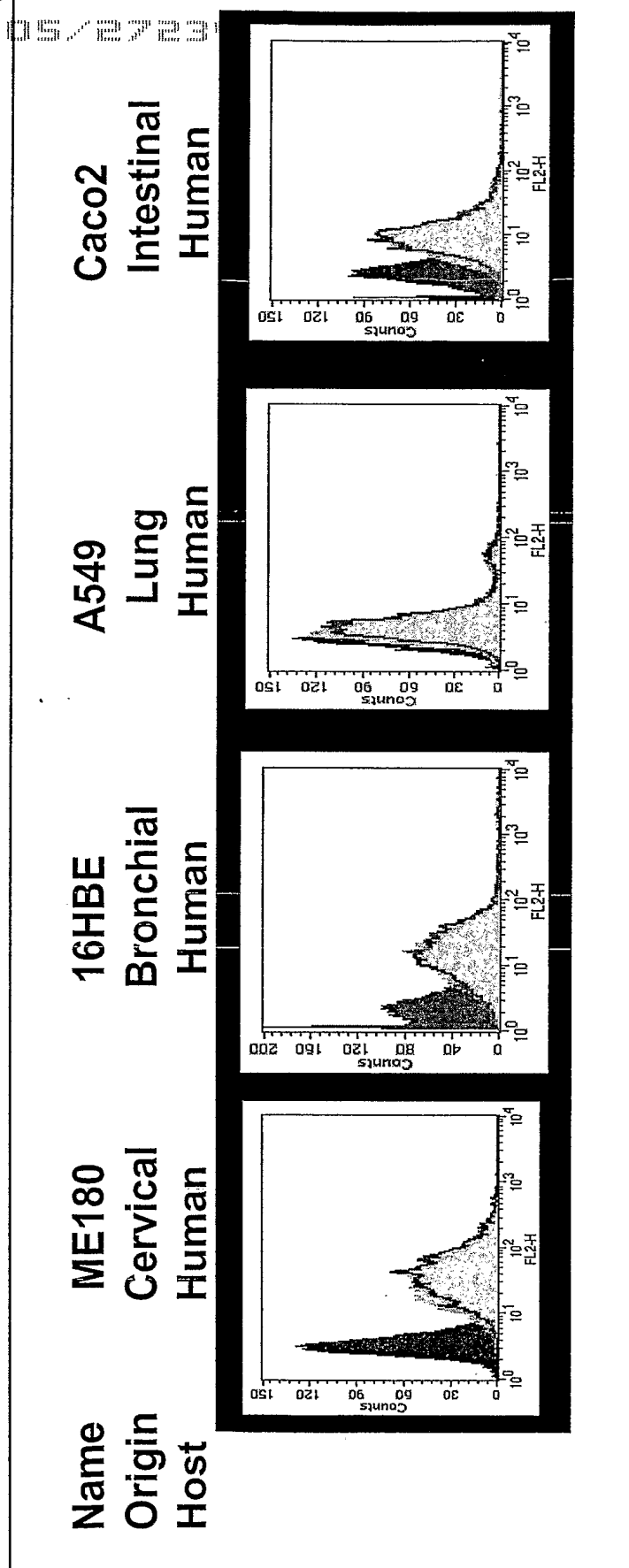
439/487

AI-4										
		aa	M1	M3	M5	M18	M49	M6	M12	
M12										
19224134	LPXTG	698	gas15 44%in297aa	M3-0098 49%in254aa				M6-0157 74%in703aa		protein F
19224135	VPXTG	756	gas15 54%in747aa	M3-0098 55%in751aa	orf78 80%in484aa	M18-0126 59%in483aa		M6-0157 51%in275aa		Cpa
19224137	QVXTG	342	gas16 40%in354aa	M3-0100 61%in344aa	orf80 65%in384aa	M18-0128 62%in344aa				EflSLA (fimbrial)
19224139	LPXAG	189	gas18 31%in206aa	M3-0102 99%in183aa	orf82 98%in189aa	M18-130 97%in189aa				Orf2
19224141	LPXTG	1161		M3-0104 59%in612aa	orf84 50%in701aa	M18-0132 50%in701aa				protein F2

Figure 201

Figure 202

GBS80 recombinant protein does not bind to epithelial cells

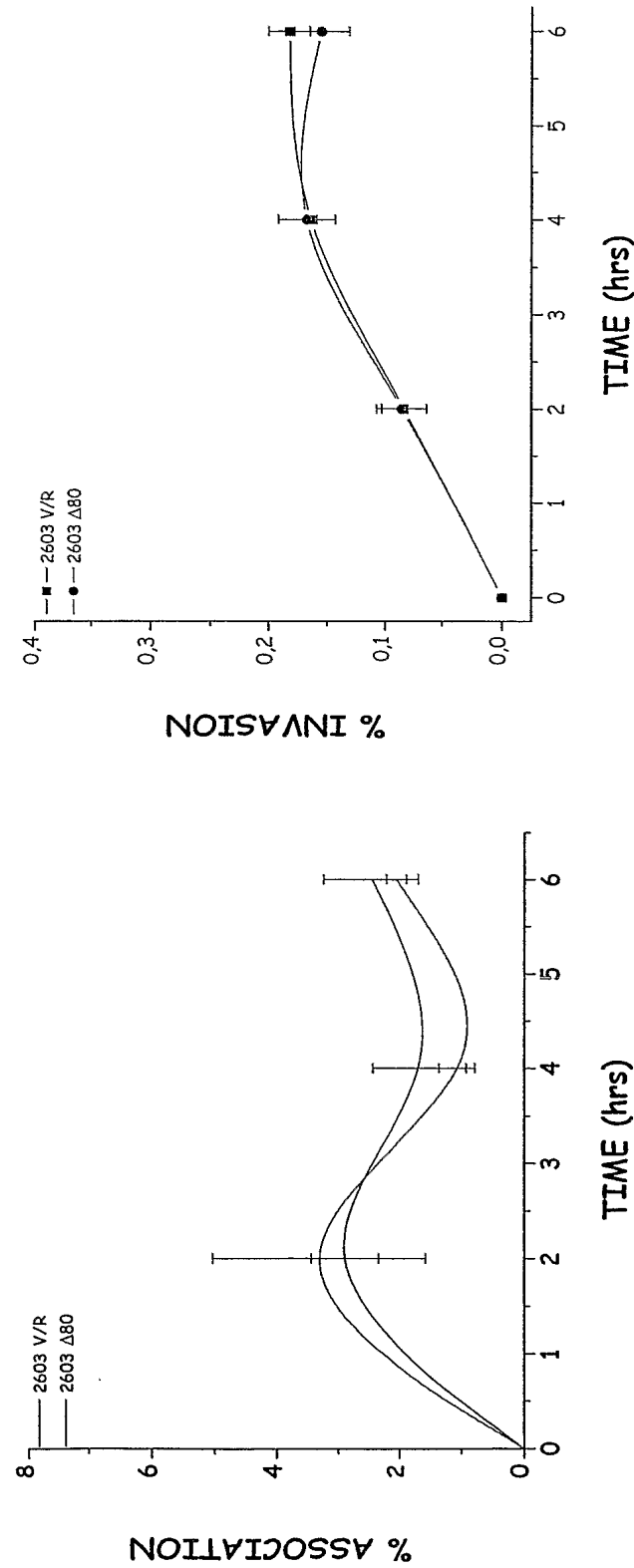


Epithelial cells were incubated in the presence or absence of GBS80 protein and then a mouse a-GBS80 polyclonal antibody added. The cell were then stained with FITC-conjugated a-mouse IgG antibody. The violet area indicates cells treated with FITC-conjugated antibody alone. GBS80 binding, expressed as Dmean channel values, was measured by FACSscan cytometer as difference in fluorescence intensity between cell incubated with or without GBS80. The same protocol was used for GBS101 protein binding to epithelial cells

441/487

Figure 203

Deletion of GBS80 protein does not affect the ability of GBS to adhere and invade ME180



ME180 cervical carcinoma epithelial cells were infected with GBS 2603 wild type or 2603 D80 isogenic mutant. After 2h infection, non-adherent bacteria were washed off and infection prolonged for further 2h and 4h. In invasion experiments, after each time point followed a 2h antibiotic treatment. Cells were then lysed with 1% saponin and lysates plated on TSA plates.

Figure 204

GBS80 binds to ECM proteins

ELISA with purified ECM components and native GBS80 protein

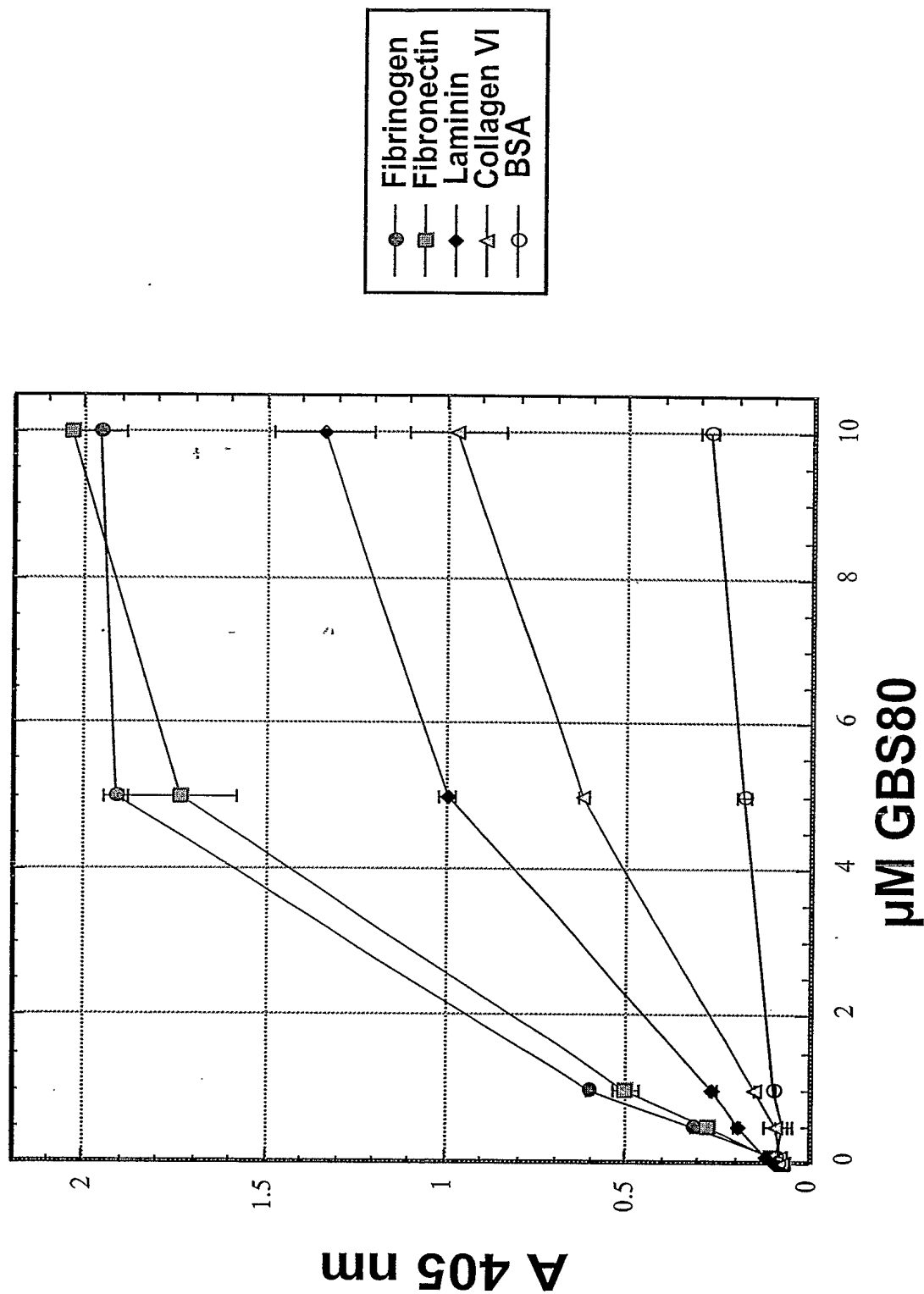
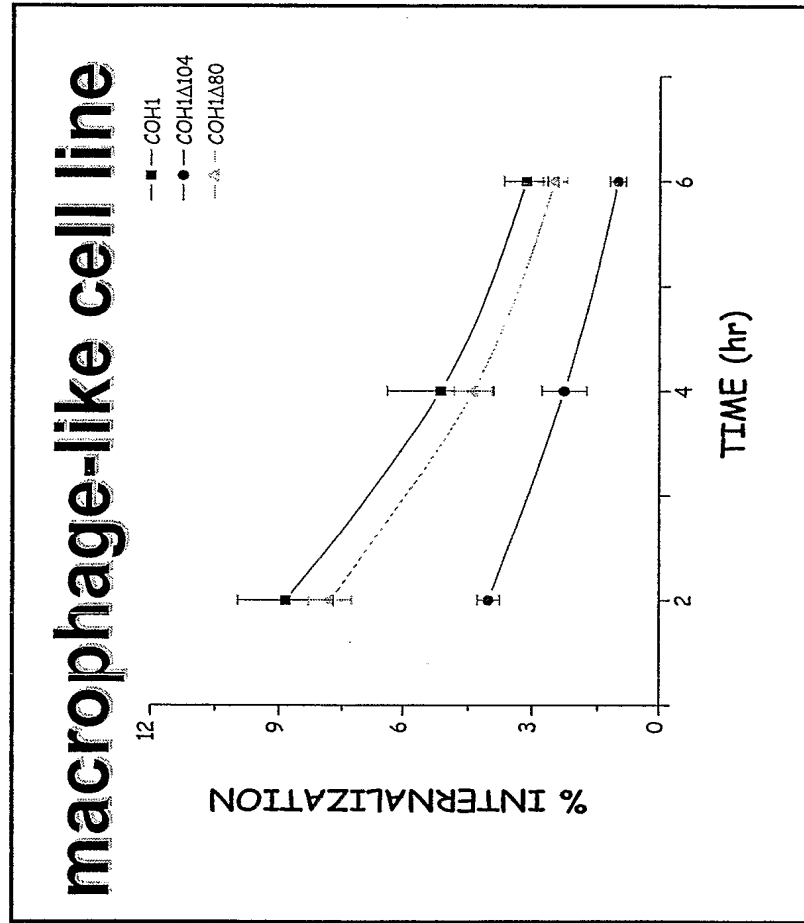


Figure 205

Deletion of GBS104 protein, but not GBS80, reduces the capacity of GBS to invade J774



J774 cells were infected with GBS COH1 wild type or COH1ΔGBS104/COH1ΔGBS80 isogenic mutants. After 1h infection, non-adherent bacteria were washed off and intracellular bacteria recovered at 2h, 4h and 6h post-antibiotic treatment. At each time point cells were lysed with 0.25% Triton X-100 and

Figure 206

**GBS104 knockout mutant strain translocates
through an epithelial monolayer less efficiently than
the isogenic wild type**

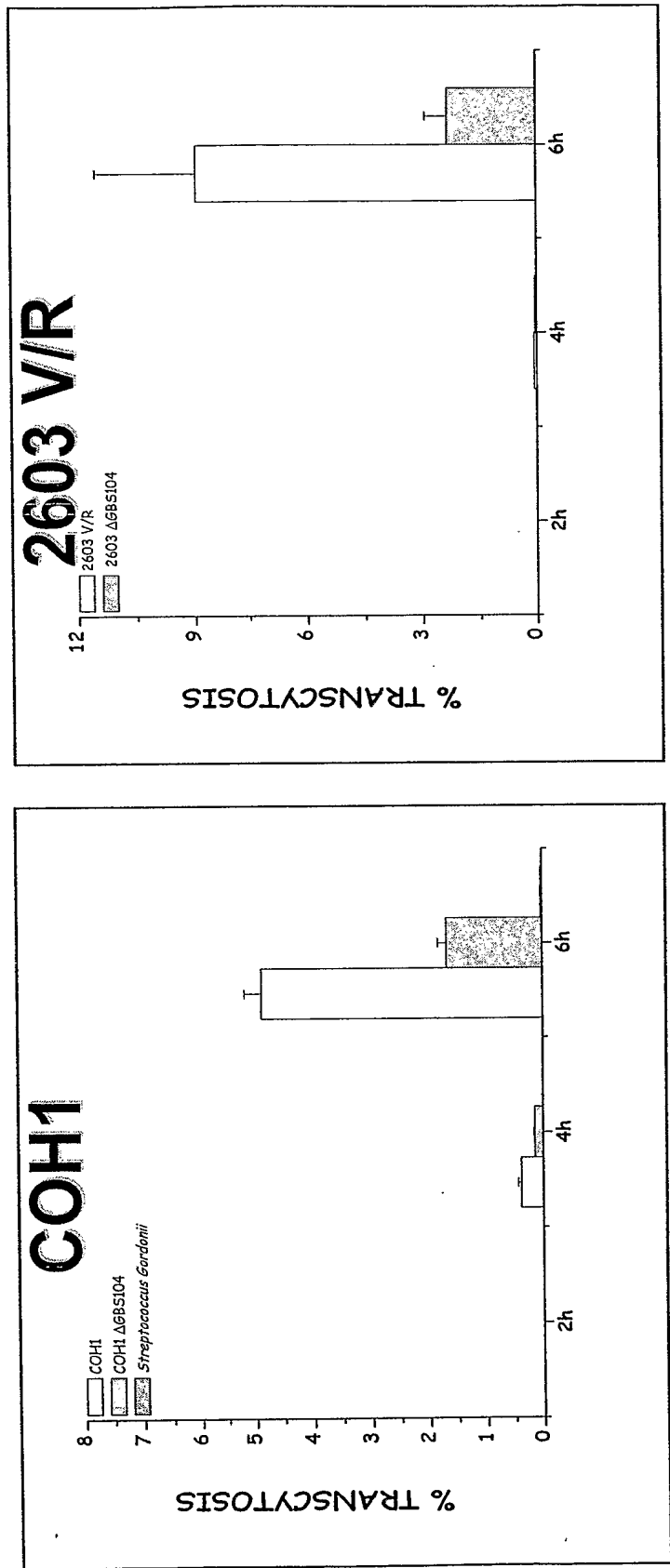
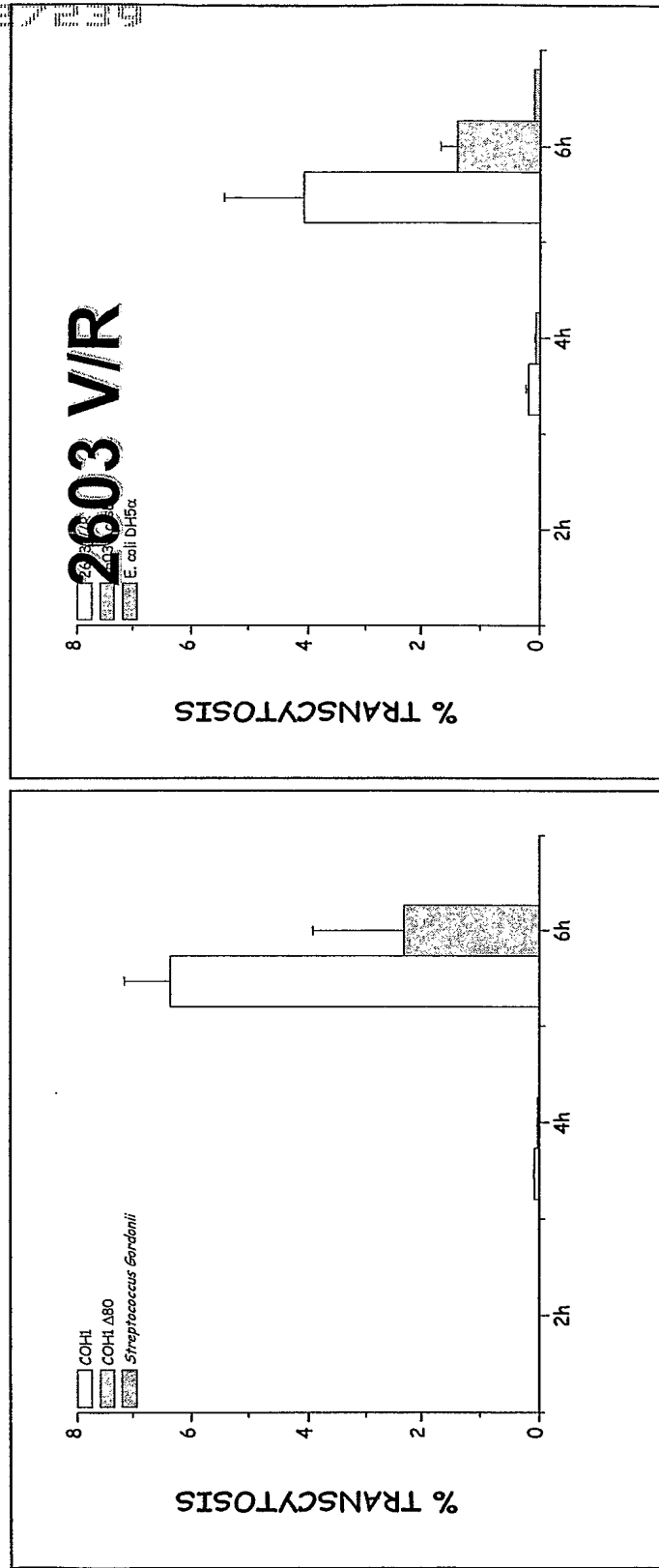


Figure 207

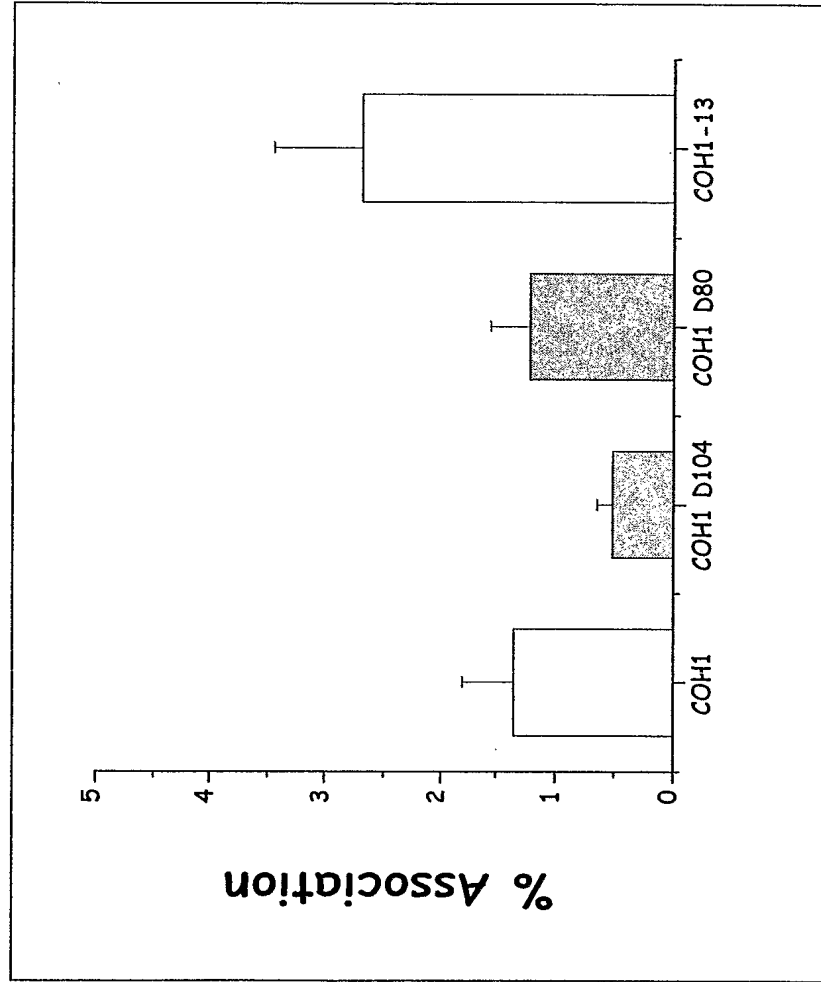
GBS80 knockout mutant strain partially loses the ability to translocate through an epithelial monolayer



Epithelial cells monolayers were inoculated with each bacterium in the apical chamber of a transwell system for 2h and then non-adherent bacteria washed off. Infection was prolonged for further 2h and 4h. Samples were taken from the media of the basolateral side and the number of colony forming units measured. Transepithelial electrical resistance measured prior and after infection gave comparable values, indicating the maintenance of the integrity of the monolayer.

Figure 208

GBS adherence to HUVEC endothelial cells



HUVEC cells were infected with GBS COH1 wild type or COH1DGBS104/COH1DGBS80 isogenic mutants. After 1h infection, non-adherent bacteria were washed off and cells lysed with 1% saponin and lysates plated on TSA plates.

COH1 strain growth rate

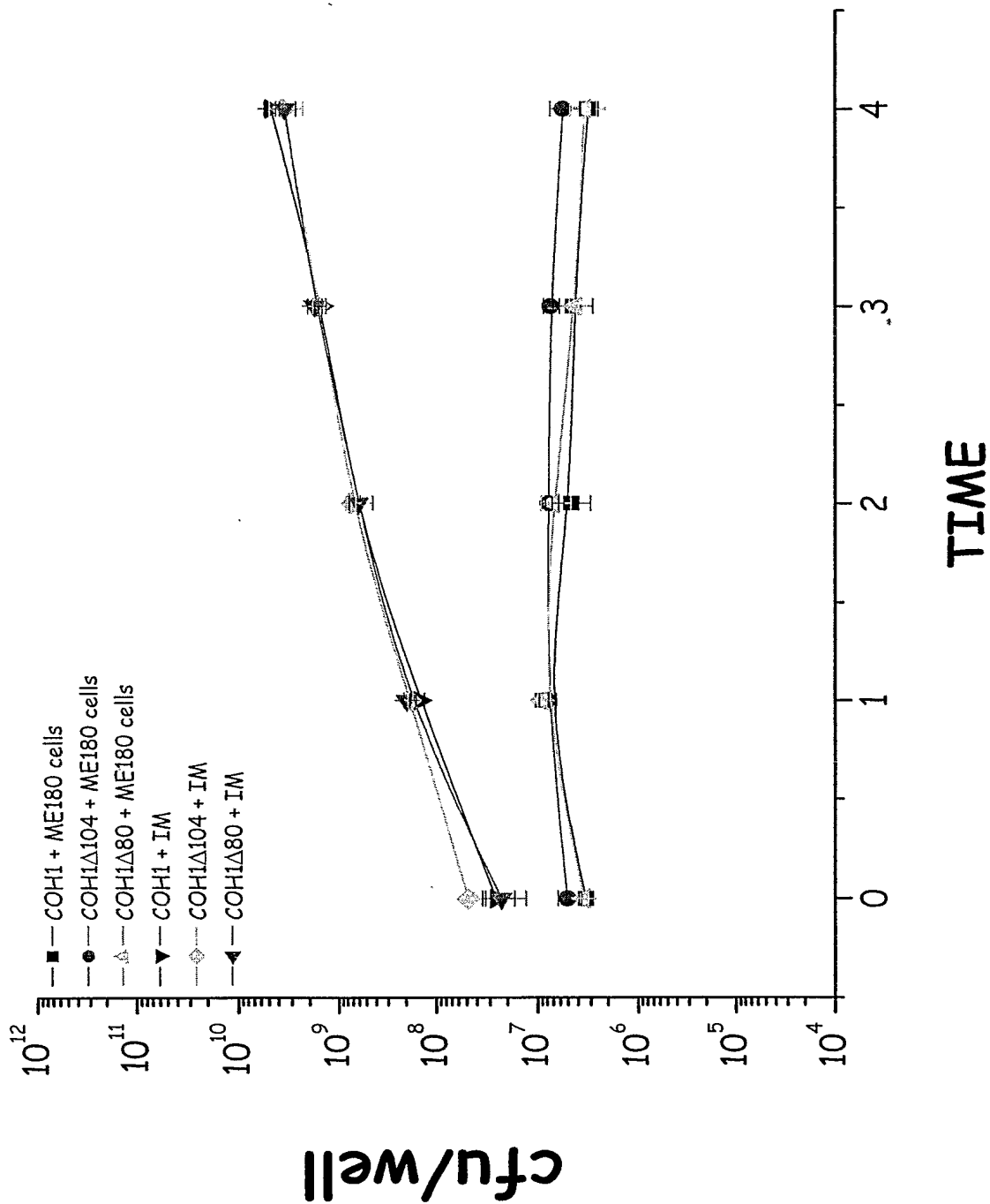


Figure 210

Binding of recombinant GBS104 protein to epithelial cells

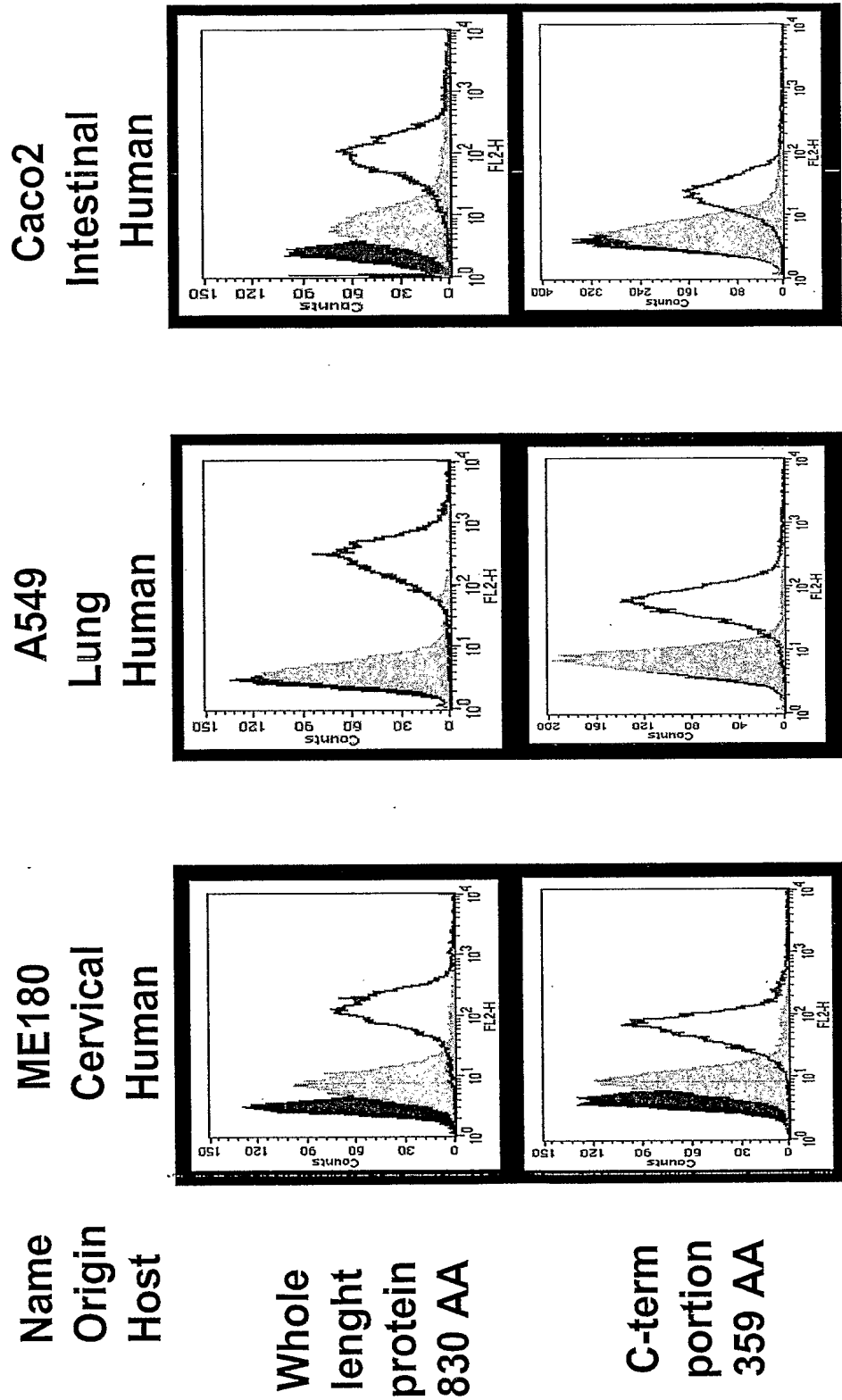
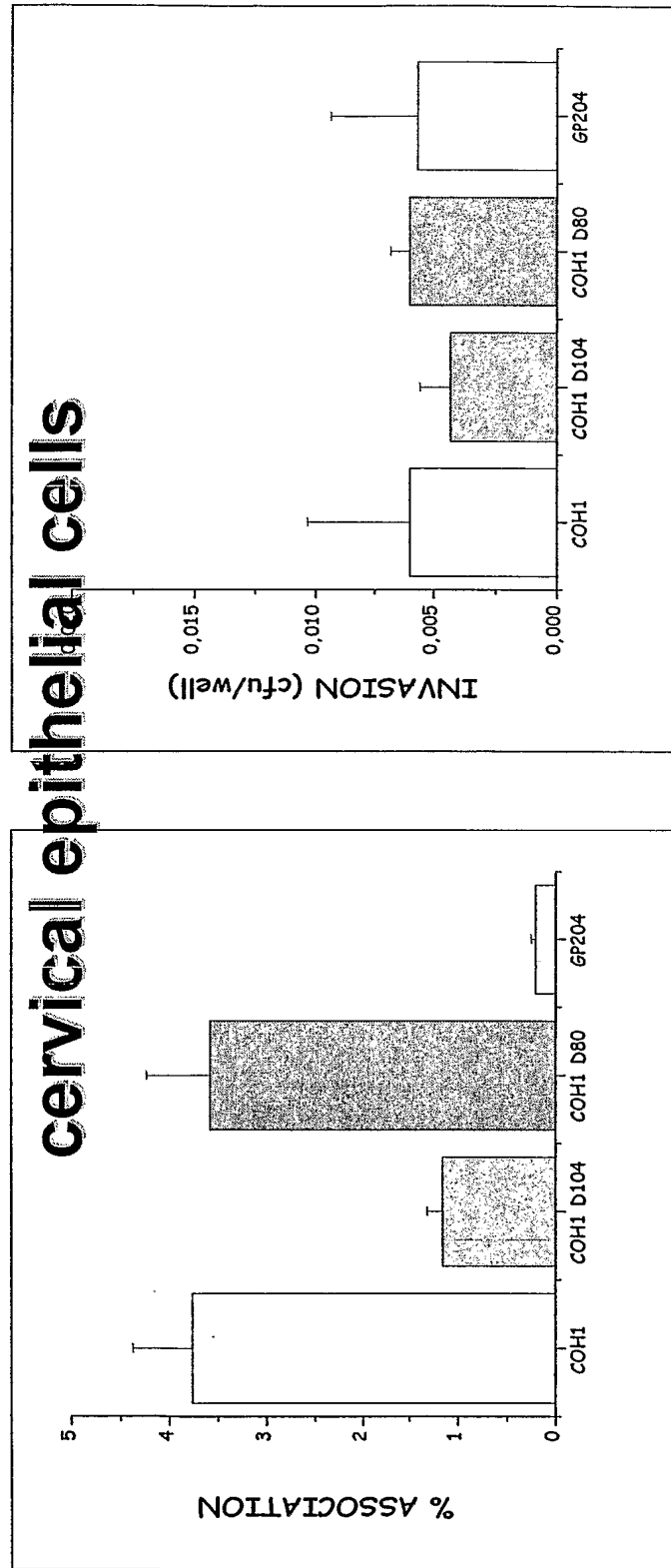


Figure 211

Deletion of GBS104 protein in the GBS strain COH1 reduces the ability of GBS to adhere to ME180

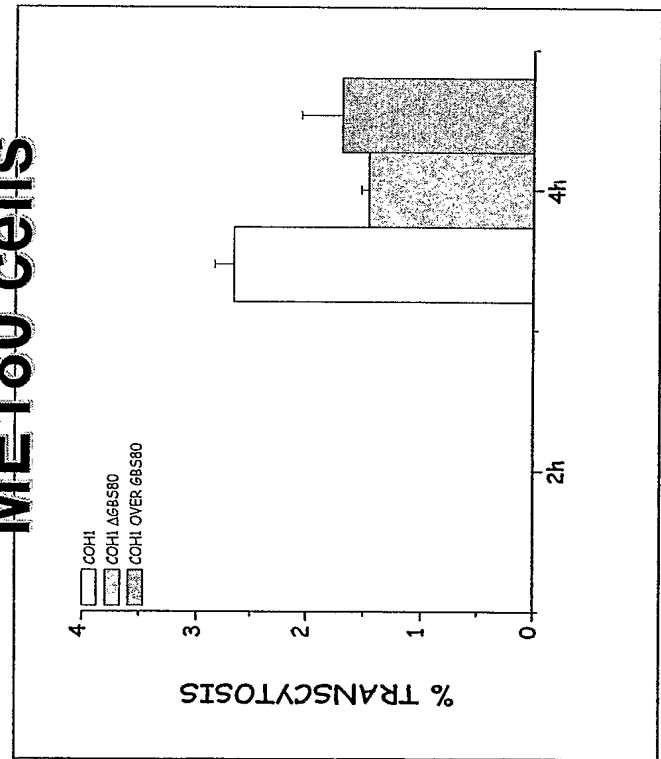


ME180 cervical carcinoma epithelial cells were infected with GBS COH1 wild type or COH1DGBS104/ COH1DGBS80 isogenic mutants. After 1h infection, non-adherent bacteria were washed off and cells lysed with 1% saponin and live bacteria plated on TSA plates

Figure 212

**COH1 overexpressing GBS80 protein has
an impaired capacity to translocate
through an epithelial monolayer**

ME180 cells



Caco2 cells

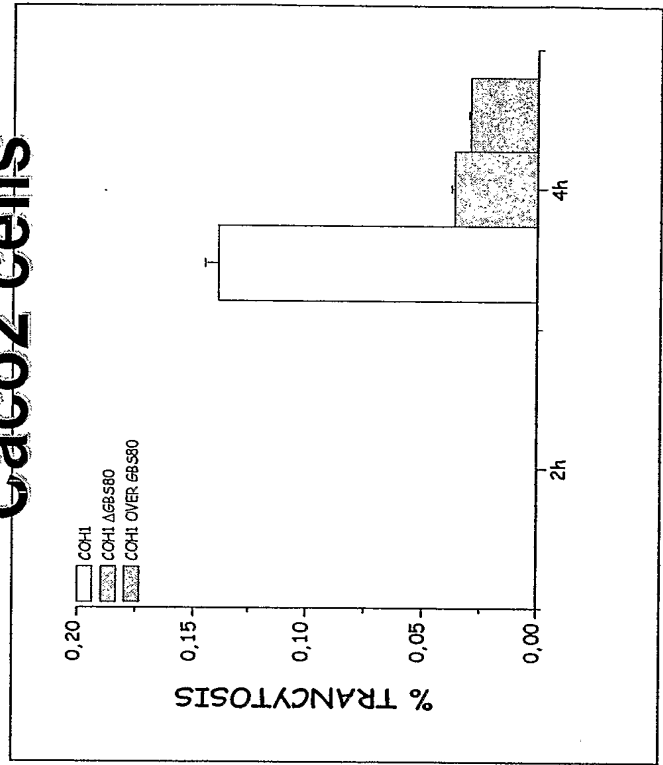


Figure 213

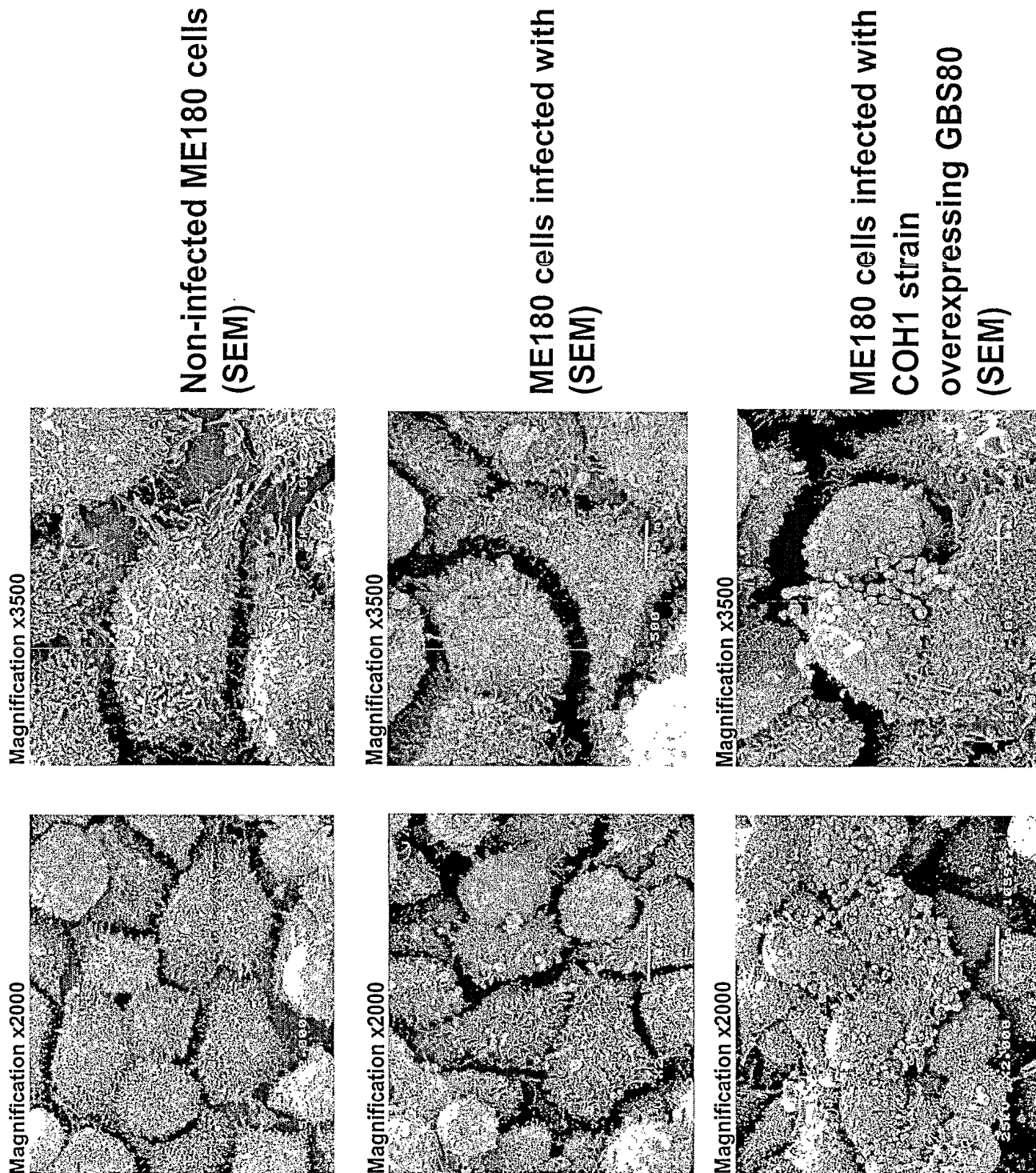


Figure 214

OH1 infection of ME180 cells

F-actin Blue

α -serotype III capsule Red

α -GBS80 Green

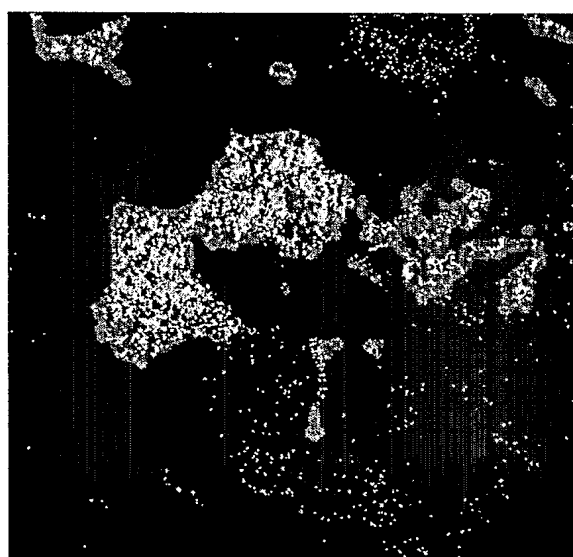
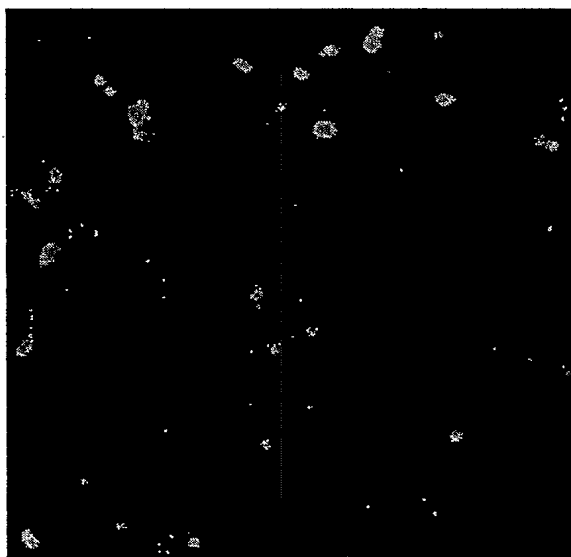
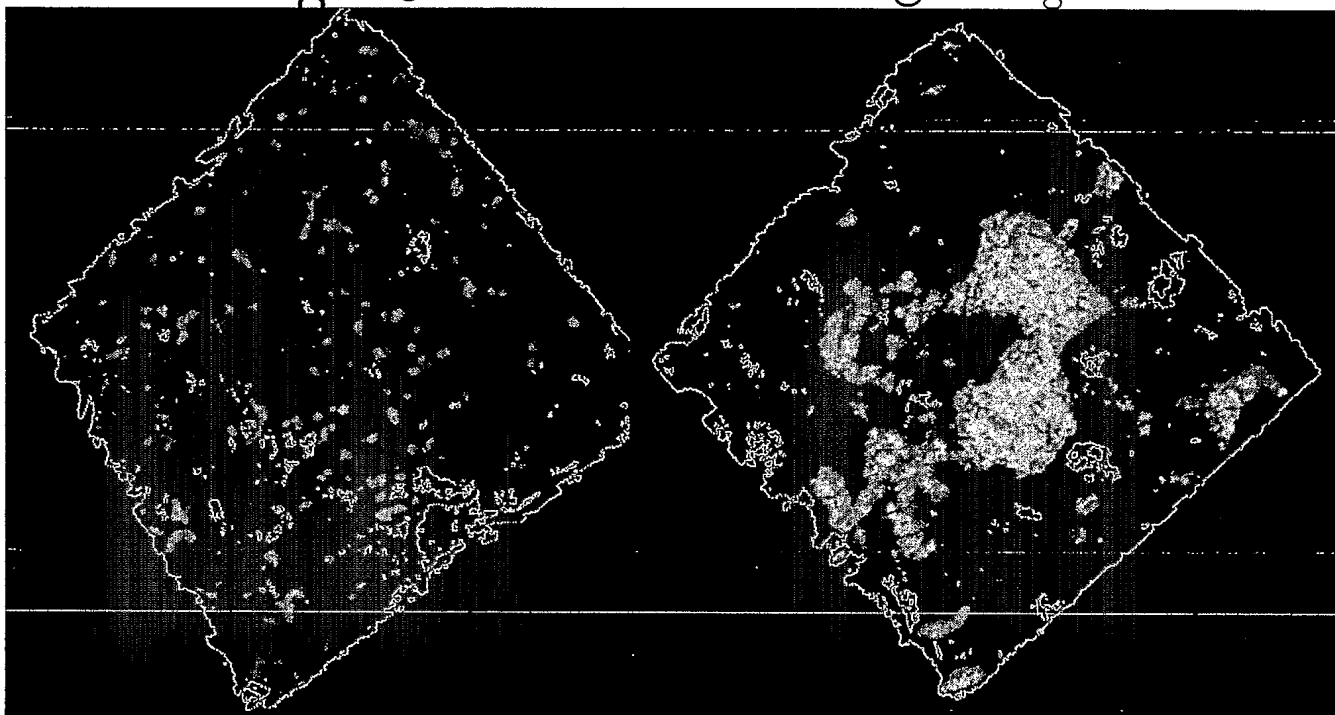
OH1 overexpressing GBS80

infection of ME180 cells

F-actin Blue

α -serotype III capsule Red

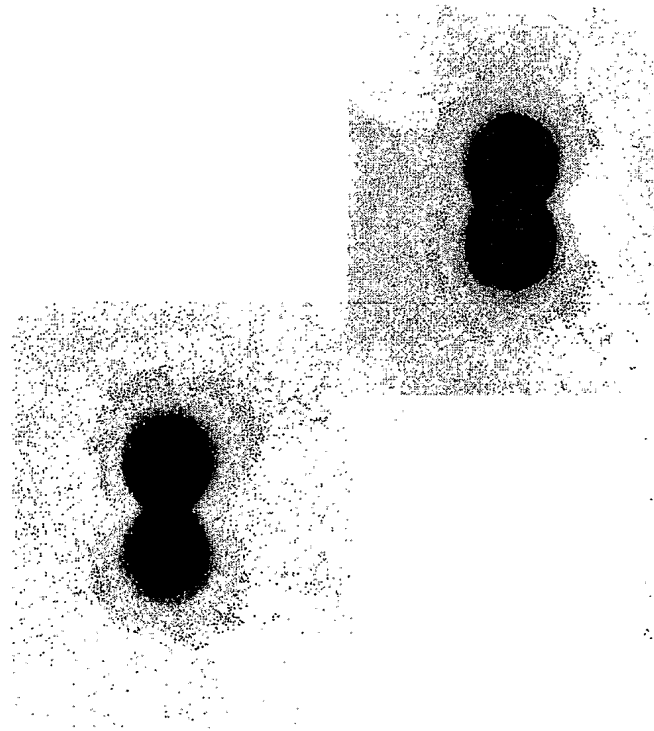
α -GBS80 Green



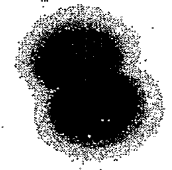
453/487

Figure 215

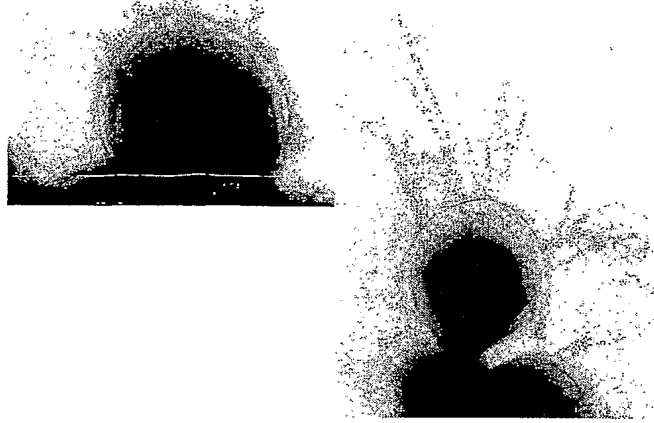
515 WT



515 Δ59



515 Δ59p59



$\alpha 59$

$\alpha 59$

Figure 216

WO 2006/078318

PCT/US2005/027239

454/487

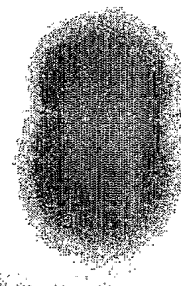
PCT/US2005/027239

515 Δ 67p67

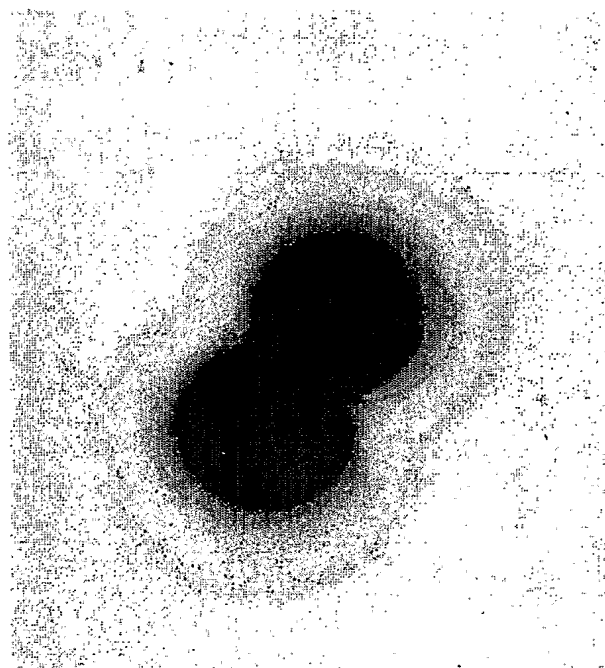


α 67

515 Δ 67



515 WT



α 67

Figure 217
GBS 67 binds to fibronectin

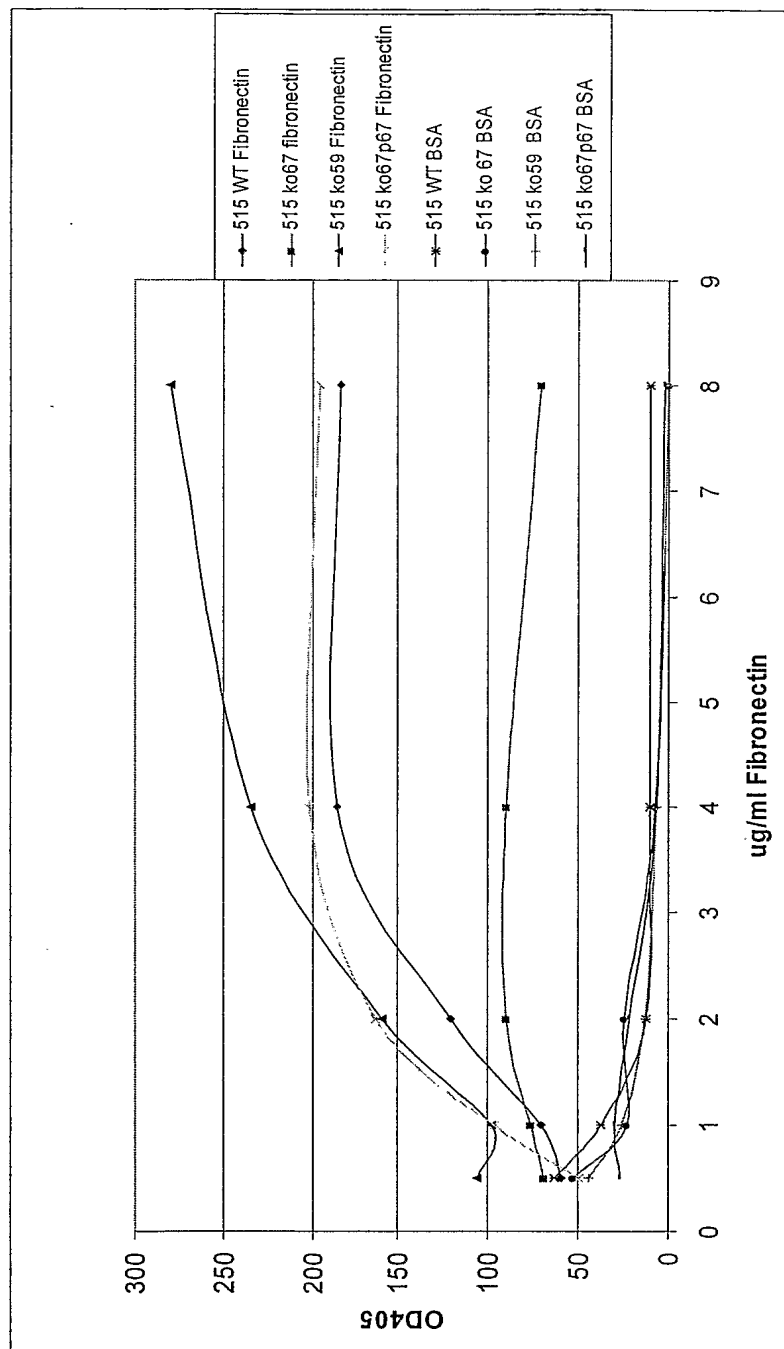


Figure 218

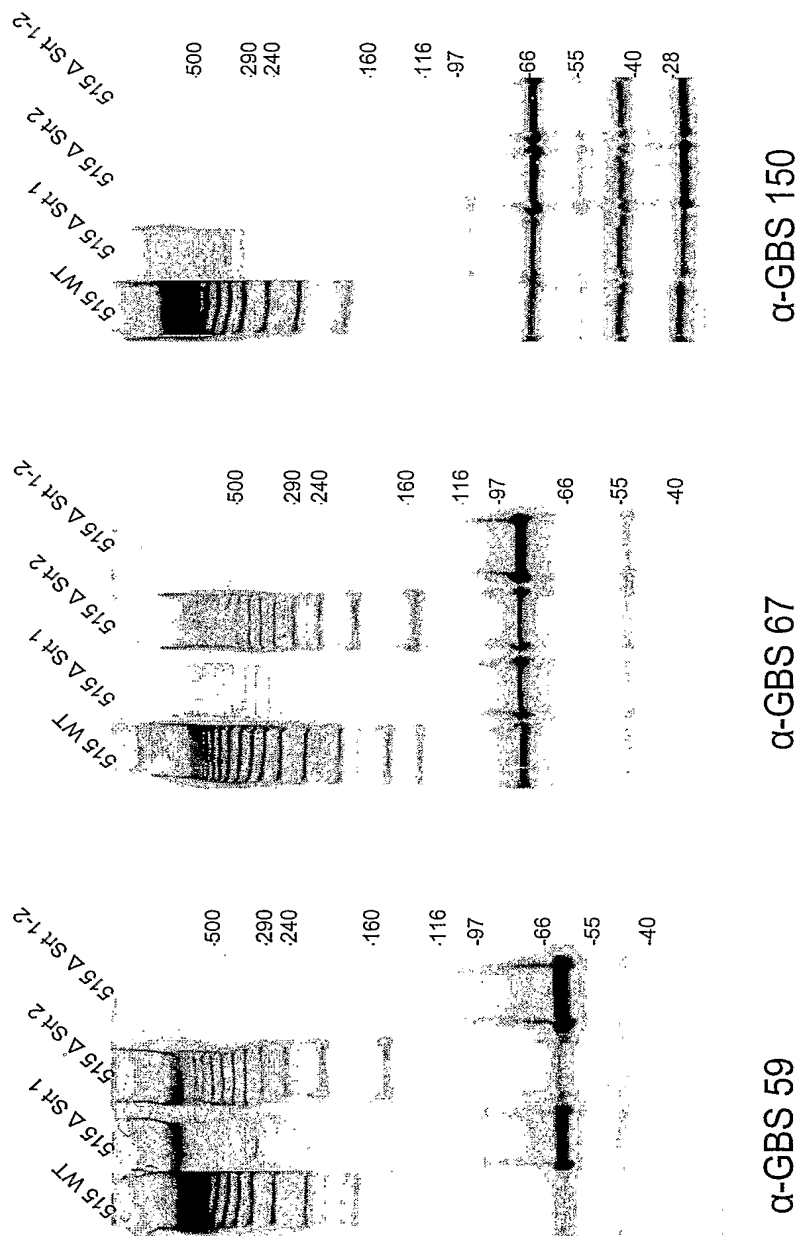
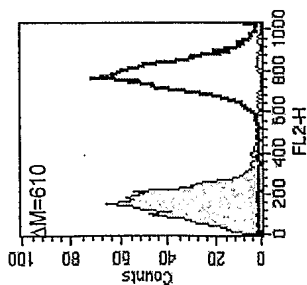
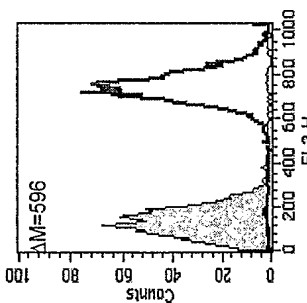


Figure 219

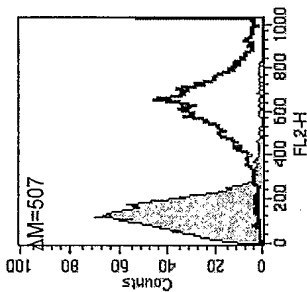
515 WT



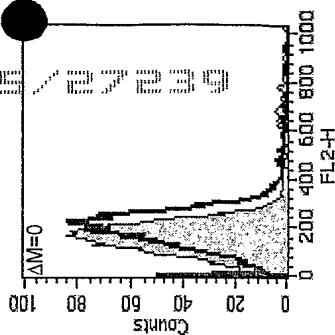
515 Δ Srt 1



515 Δ Srt 2

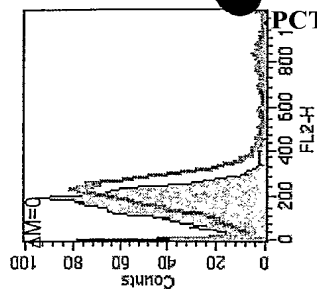
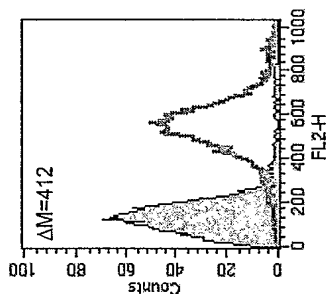
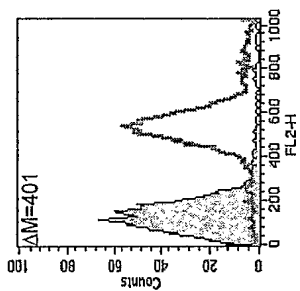
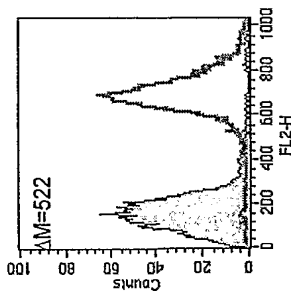


515 Δ Srt 1.2



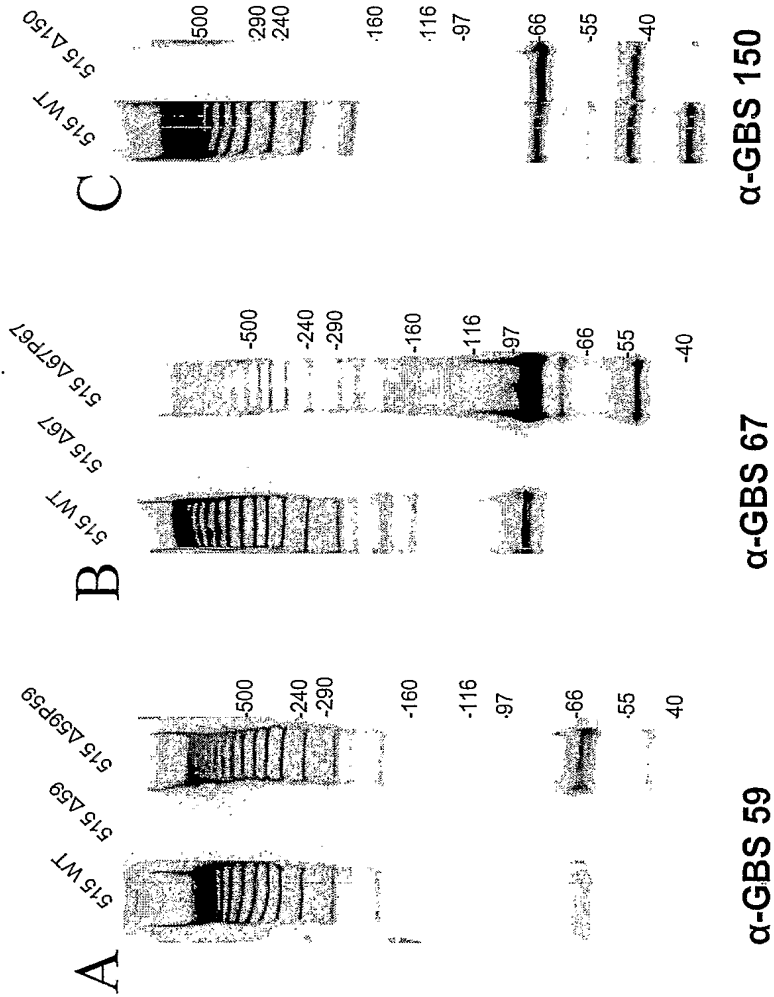
457/487

α67



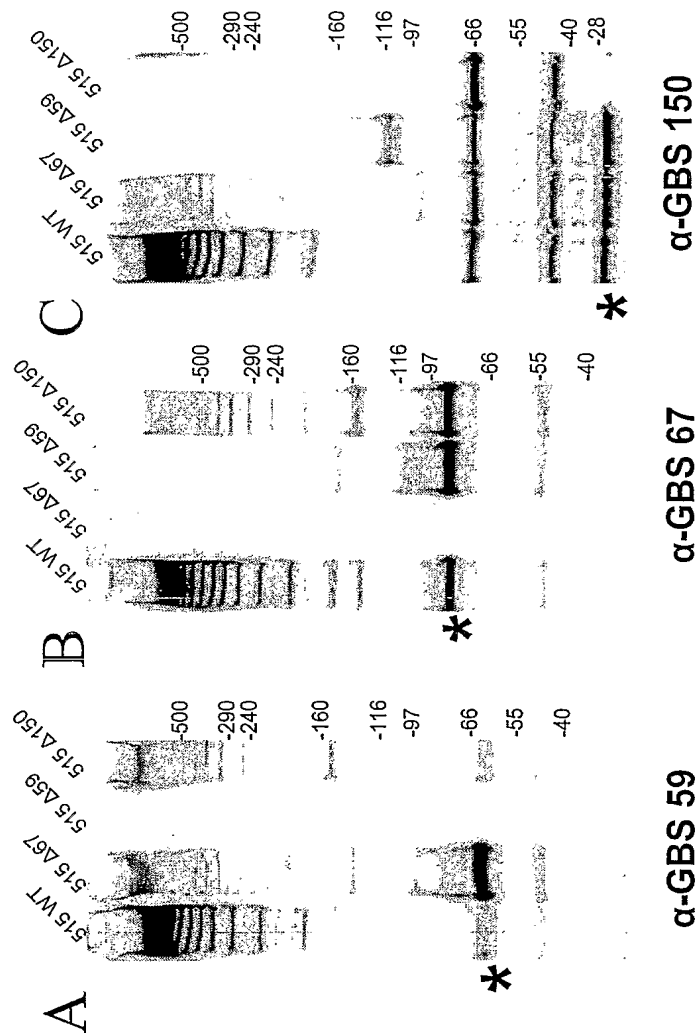
α59

Figure 220



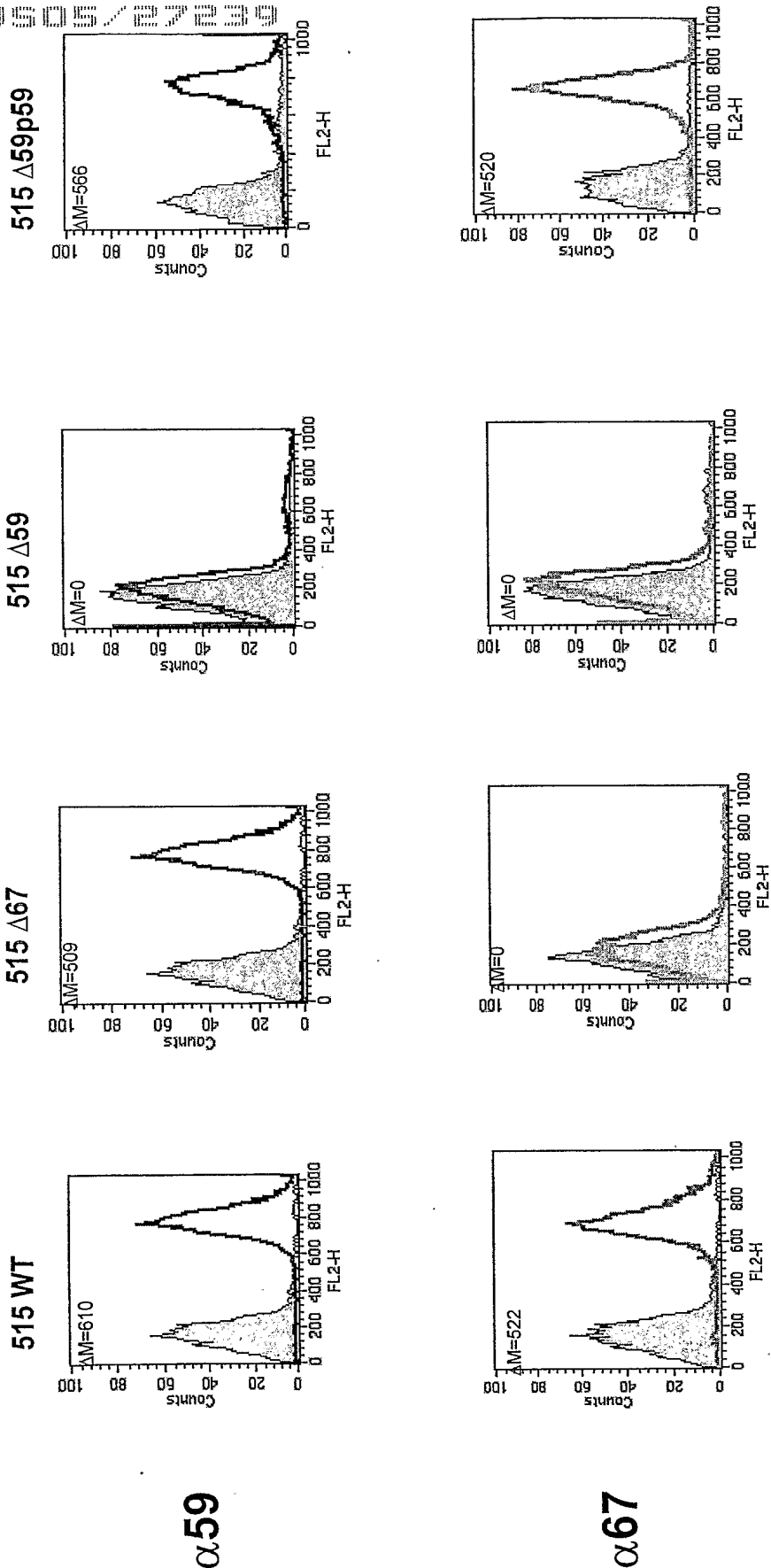
PCT/US05/27239 459/487

Figure 221



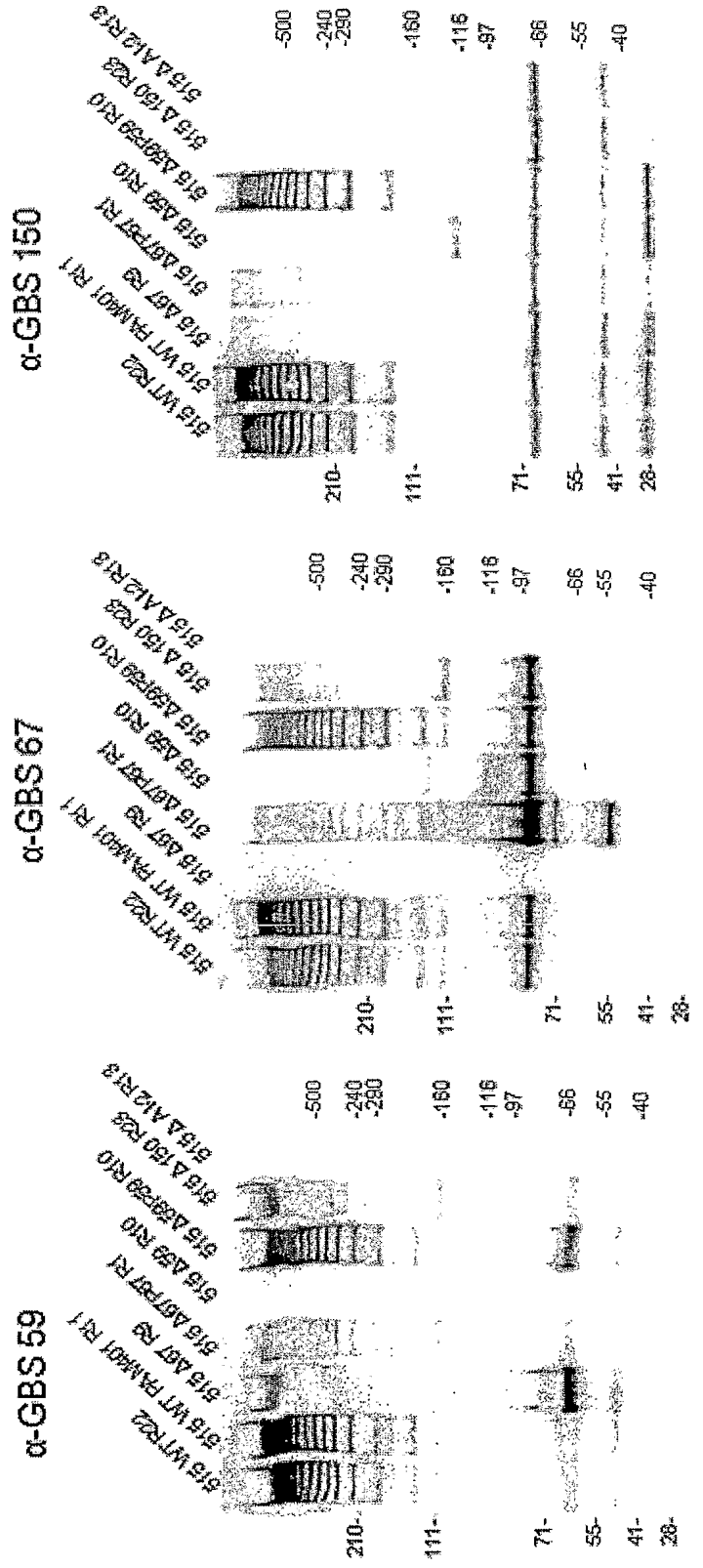
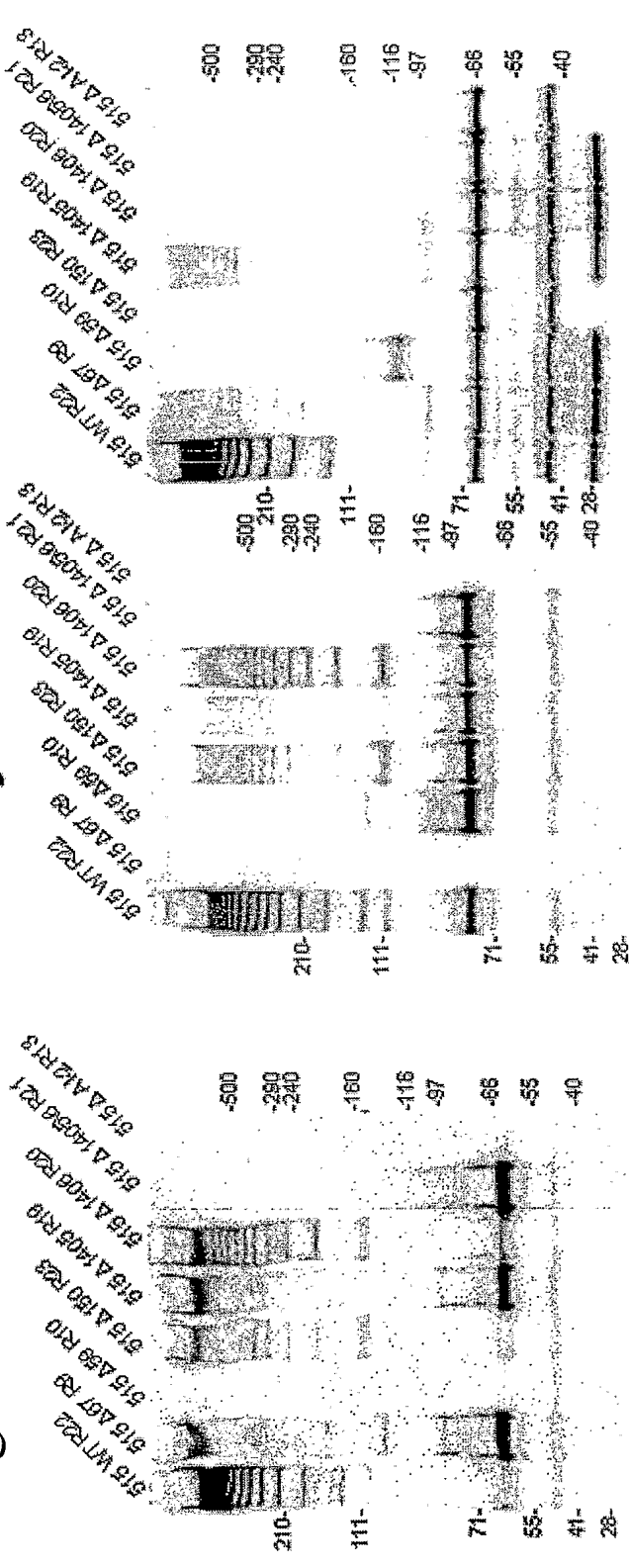
PCT/US05/27239

Figure 222



Summary WB

Figure 223



GBS strain % AA identity

7357b (Ib)	100
5518 (Ib)	100
5364 (V)	100
1999 (IV)	100
5408 (VIII)	98
coh31 (III)	98
d136c (III)	98
nem316 (III)	98

dk1 (Ia)	100
dk8 (Ia)	100
davis (Ia)	100
5551 (Ia)	100
2986 (Ia)	100
2110 (V)	100
2210 (IV)	100

18RS21 (II)	100
3050 (II)	100
2141 (II)	100
1998 (III)	100
2928 (VII)	99,9

2274 (IV)	99,9
2129 (Ib)	99,7
5401 (II)	99,8

GBS 59 allelic variants

cjb111 (V)
674 aa

515 (Ia)
675 aa

2603 (V)
705 aa

H36b (Ib)
693 aa

75%

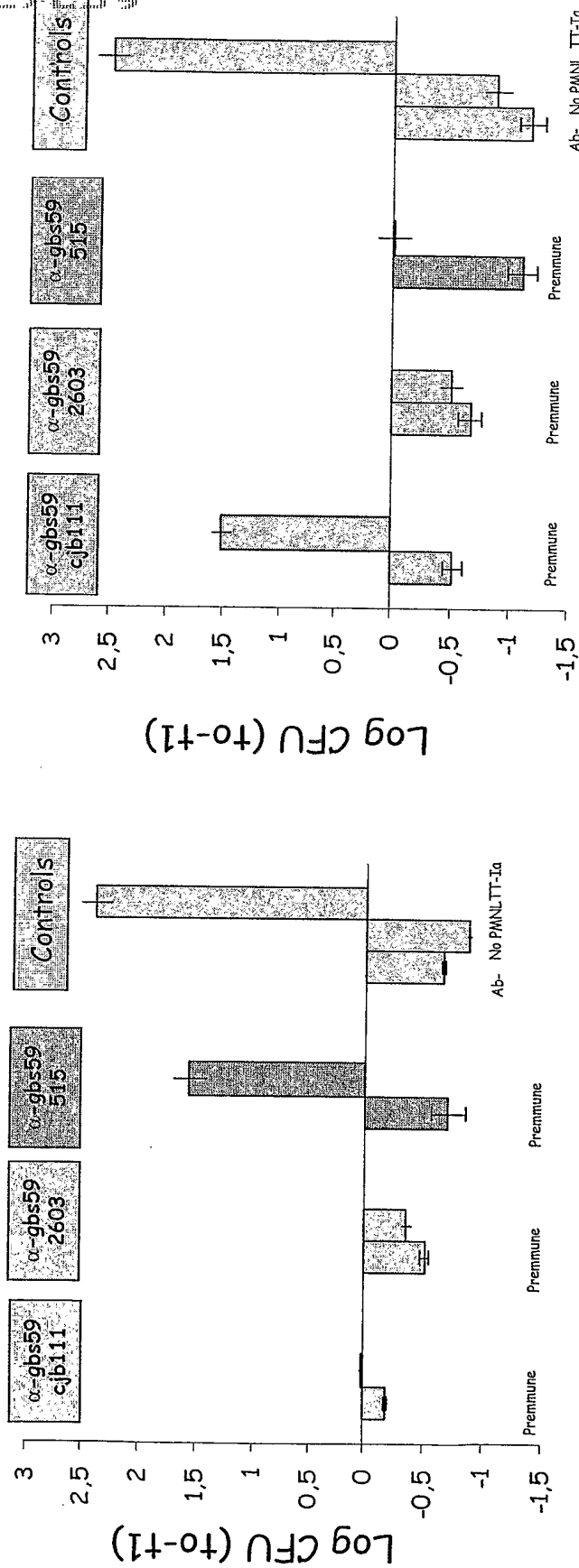
48%

65%

Figure 224

Figure 225

GBS 59 is opsonic only against homologous strain



• 515 (Ia) GBS strain

• cjb111 (V) GBS strain

Figure 226 A

		GBS 59		
GBS strains	Type	PCR	FACS (a-cjb111)	FACS (a-2603)
DK1	Ia	+	565	
DK8		+	559	
Davis		+	577	
515		+	583	0
090		+	0	0
2986		+	443	
5551		+	524	
H36B	Ib	+	0	410
7357b-		+	596	
5518		+	190	
D136C	III	+	504	
COH31		+	505	
1998		+	59	510
18RS21	II	+	0	353
DK21		+	249	0
3050		+	0	570
5401		+	0	400
2141		+	0	371
CJB111	V	+	625	0
2603		+	0	73
5364		+	593	
2110		+	590	0
2274	IV	+	0	400
1999		+	594	
2210		+	636	
5408	VIII	+	537	
CJB110	NT	+	0	0
1169		+	227	0

		GBS 59		
GBS strains	Type	PCR	FACS (a-cjb111)	FACS (a-2603)
A909	Ia	-	22	0
2177	Ib	-	75	
COH1	III	-	0	
M732		-	0	
M781		-	17	
5376		-	60	
5435		-	55	
SMU071	VIII	-	0	0
JM9130013		-	0	

Figure 226 B

Figure 227 A

		FACS (D Mean)				
GBS strains	Type	GBS 80	GBS 104	GBS 67	GBS 322	GBS 59
DK1	Ia	0	0	478	153	565
DK8		0	0	475	213	559
Davis		0	0	430	86	577
515		0	0	409	227	583
090		0	0	0	0	0
A909		46	29	0	0	0
2986		0	0	397	0	443
5551		0	0	485	36	524
2177	Ib	477	355	66	323	0
H36B		0	0	444	105	410
7357b-		91	0	316	102	596
5518		31	0	162	0	190
COH1	III	305	226	0	130	0
D136C		40	40	406	460	504
COH31		0	0	273	479	505
M732		141	101	0	292	0
M781		111	136	0	224	0
1998		140	77	350	288	510
5376		165	156	0	76	0
5435		93	100	0	88	0
18RS21	II	0	0	103	471	353
DK21		0	0	331	342	249
3050		71	46	460	188	570
5401		75	28	618	135	400
2141		0	0	370	76	371
CJB111	V	365	236	481	58	625
2603		62	0	105	293	73
5364		454	281	394	463	593

2110		0	0	589	0	590
2274		123	62	484	161	400
1999	IV	0	389	453	55	594
2210		0	0	574	0	636
SMU071		556	393	74	170	0
JM9130013	VIII	587	436	72	133	0
5408		0	0	433	0	537
CJB110		0	0	245	587	0
1169	NT	0	0	443	213	227
D Mean > 200		6/37 (16%)	7/37 (19%)	24/37 (65%)	14/37 (38%)	24/37 (65%)

Figure 227B

Figure 228

		FACS (ΔMean)																Δmean
GBS Strain	Type	GBS 80		GBS 104		GBS 322		GBS 67		GBS 67		GBS 59		GBS 59		GBS 59		neg. control
		142-F		Mab		86		81		H36B		2603		CJB111		515		
cdc-1	II	114	95	0	0	122	122	360	341	422	403	92	73	254	235	306	287	19
cdc-2	IB	173	69	92	0	95	75	552	448	590	486	135	31	635	531	197	93	104
cdc-3	II	566	508	360	302	85	60	364	306	433	375	111	53	448	390	310	252	58
cdc-4	V	524	432	337	245	284	204	577	485	625	533	105	13	674	582	303	211	92
cdc-5	II	140	0	0	0	462	300	487	297	563	373	175	0	373	183	440	250	190
cdc-6	V	544	484	361	301	95	95	586	526	601	541	55	0	686	626	302	242	60
cdc-7	III	155	116	44	5	134	118	95	56	138	99	74	35	92	53	91	52	39
cdc-8	III	347	304	192	149	74	62	98	55	170	127	72	29	88	45	108	65	43
cdc-9	II	89	65	0	0	226	191	390	366	504	480	181	157	317	293	410	386	24
cdc-10	IA	46	24	0	0	152	152	494	472	531	509	43	21	16	0	48	26	22
cdc-11	IA	17	0	0	0	295	135	569	550	569	550	47	28	467	448	648	629	19
cdc-12	V	439	430	290	281	60	30	174	165	227	218	52	43	139	130	207	198	9
cdc-13	IA	33	0	0	0	216	146	469	436	469	436	100	67	361	328	571	538	33
cdc-14	III	78	68	10	0	213	191	50	40	85	75	38	28	69	59	67	57	10
cdc-15	III	119	53	24	0	108	98	48	0	127	61	89	23	105	39	100	34	66
cdc-16	V	363	335	177	149	310	270	70	42	127	99	48	20	130	102	128	100	28
cdc-17	III	160	0	163	0	408	248	377	217	410	250	441	281	359	199	167	7	160
cdc-18	III	49	28	0	0	239	218	34	13	36	15	16	0	49	28	56	35	21
cdc-19	III	182	101	0	0	361	280	310	229	312	231	384	303	220	139	0	0	81
cdc-20	V	348	304	203	159	380	336	166	122	211	167	114	70	232	188	128	84	44
cdc-21	II	222	132	83	0	150	60	331	241	336	246	0	0	420	330	59	0	90
cdc-22	IA	0	0	13	13	43	43	238	238	238	238	43	43	38	38	429	429	0
cdc-22 (9-6-05)		23	0	34	0	110	20	310	220	320	230	113	23	117	27	344	254	90
cdc-23	V	484	484	374	374	278	278	124	124	206	206	11	11	91	91	236	236	0
cdc-24	V	137	52	0	0	333	248	90	5	110	25	110	25	120	35	70	0	85
cdc-25	IA	0	0	0	0	351	190	530	370	565	405	495	335	442	282	625	465	160
cdc-26	II	117	2	0	0	185	70	210	95	285	170	30	0	175	60	210	95	115
cdc-27	III	323	95	34	0	498	270	346	118	406	178	424	196	314	86	64	0	228
cdc-28	V	150	92	20	0	132	74	462	404	505	447	0	0	526	468	78	20	58
cdc-29	IV	90	73	65	48	195	178	90	73	150	133	150	133	138	121	110	93	17
cdc-30	V	390	187	336	133	348	145	229	26	244	41	113	0	268	65	223	20	203
cdc-31	IA	22	0	68	0	306	182	368	244	386	262	126	2	248	124	426	302	124
cdc-32	IA	45	0	12	0	260	175	190	105	205	120	30	0	100	15	185	100	85
cdc-33	II	50	0	0	0	306	156	134	0	237	87	4	0	180	30	190	40	150
cdc-34	III	152	60	47	0	342	250	44	0	74	0	27	0	102	8	48	0	92
cdc-35	V	227	227	40	40	246	246	395	395	415	415	0	0	550	550	142	142	0
cdc-36	IB	25	15	8	0	30	20	154	144	174	164	33	23	222	212	20	10	10
cdc-37	III	168	53	61	0	361	246	82	0	133	18	83	0	132	17	75	0	115
cdc-38	II	140	14	30	0	338	212	124	0	198	72	158	32	138	12	104	0	126
cdc-39	II	126	0	0	0	316	148	466	298	514	346	438	270	184	16	34	0	168
cdc-40	V	420	366	214	160	22	0	103	49	162	108	90	36	209	155	192	138	54
cdc-41	II	146	31	15	0	380	265	330	215	425	310	140	25	280	165	315	200	115

Figure 229

Expected strain coverage

MIX GBS proteins

n. antigens FACS++	vaccine options					w/o 322			w/o 104+322			w/o 59+322		
	80+104+67+59+322	80+104+67+322	80+104+67+59	80+104+67+59	80+104+67+322	80+104+67+59	80+104+67+322	80+104+67+59	80+104+67+322	80+104+67+59	80+104+67+322	80+104+67+59	80+104+67+322	80+104+67+59
1	89%	89%	89%	80%	80%	80%	80%	80%	79%	79%	79%	79%	79%	74%
2	74%	74%	51%	71%	71%	71%	71%	64%	24%	24%	24%	24%	16%	16%
3	23%	23%	14%	17%	17%	17%	17%	16%	13%	13%	13%	13%	13%	16%

- GBS 322 but not GBS 59 is important to increase strain coverage
- GBS 59 probably could be useful to increase the vaccine strength

Assumption:

- Protein antigens that are highly accessible to antibodies confer 100% protection with suitable adjuvants

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Figure 230

GBS 59 opsonophagocytic activity is comparable to that of the four-protein mix

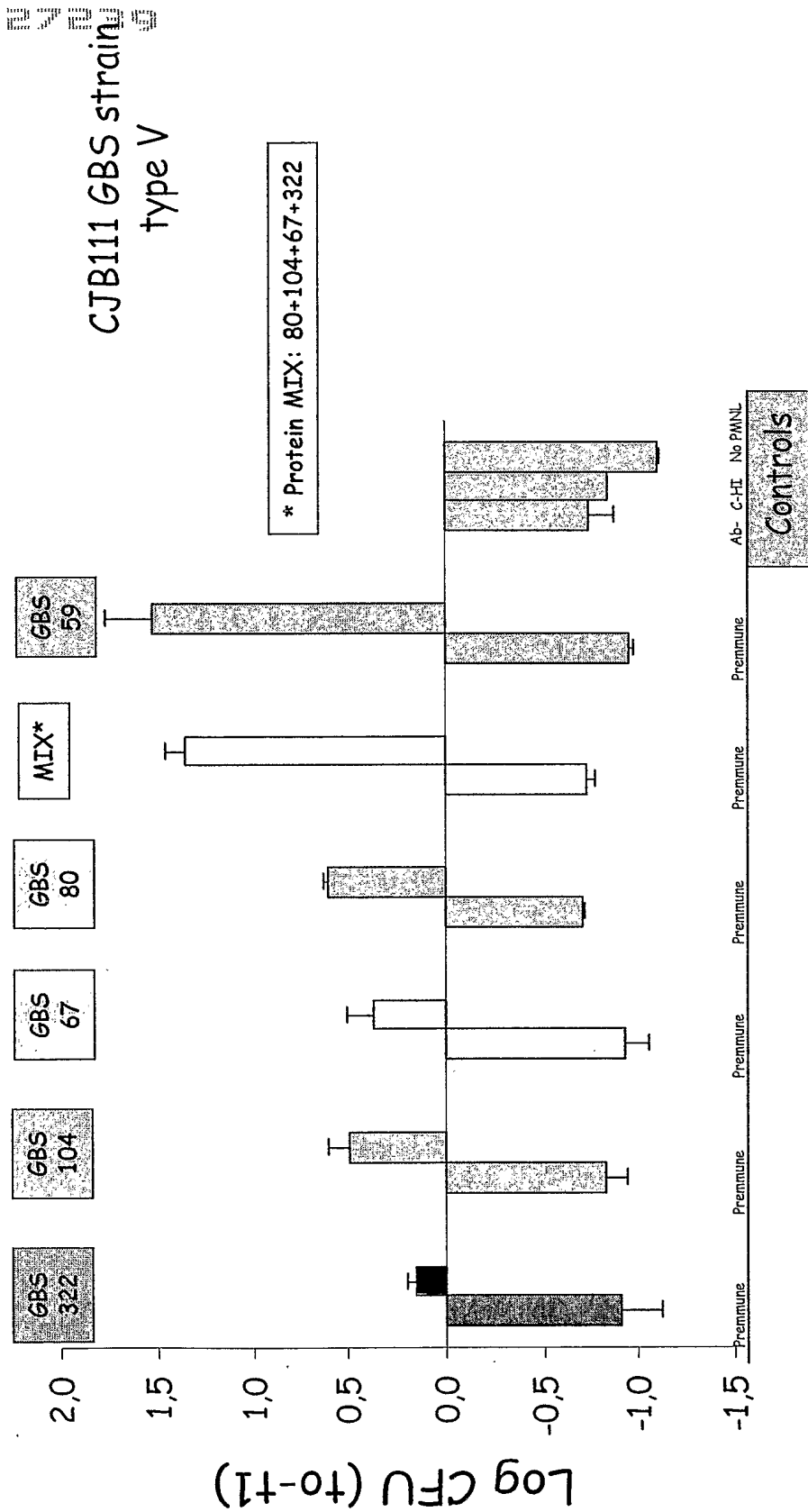


Figure 231

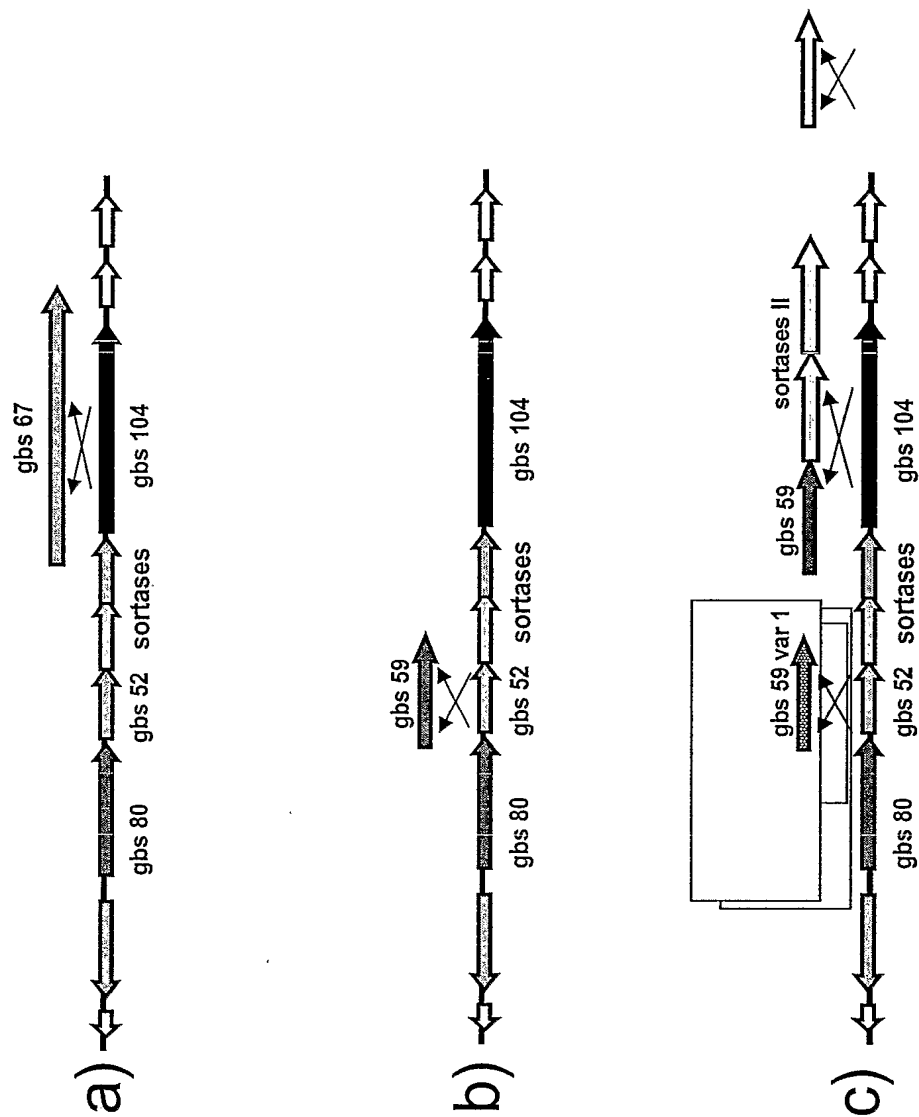
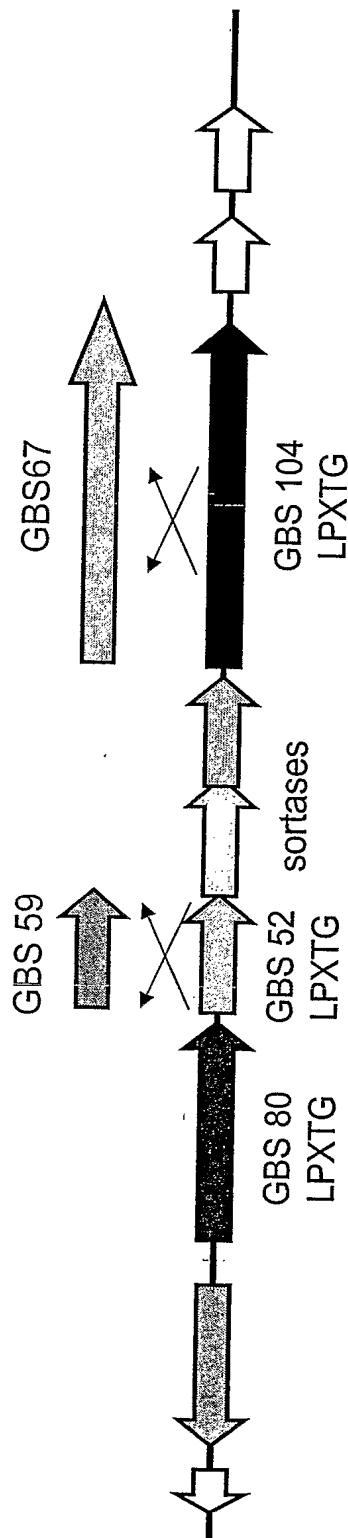


Figure 232



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Figure 233

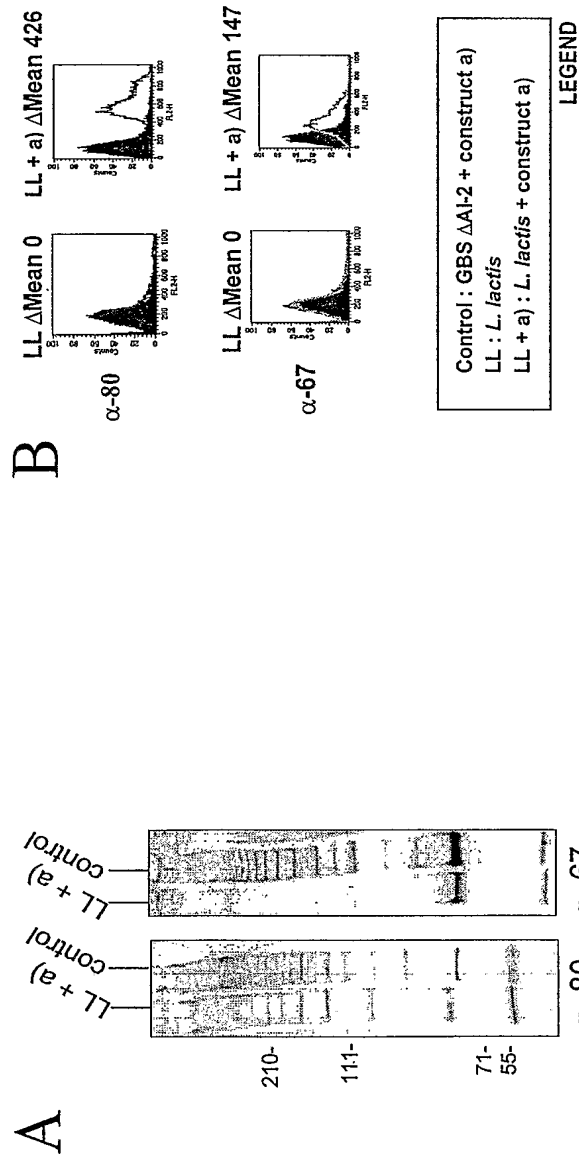
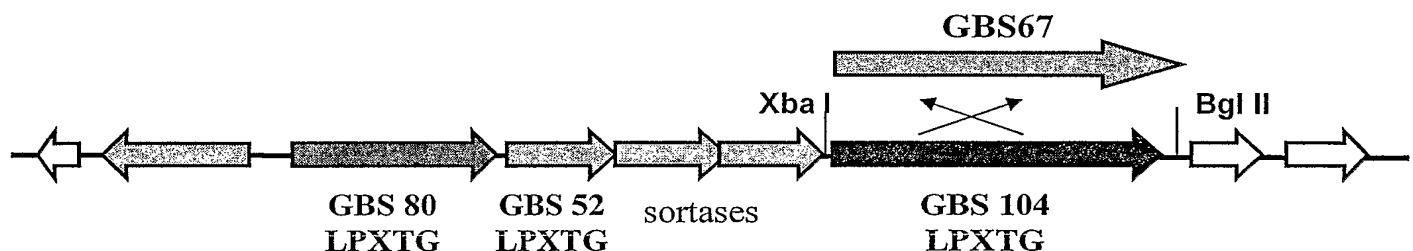


Figure 234 A

Introducing Heterologous Antigens into AI-1 pilus to Obtain Protection Across GBS Strains

1- Substitution of GBS 104 with GBS67 from Island II

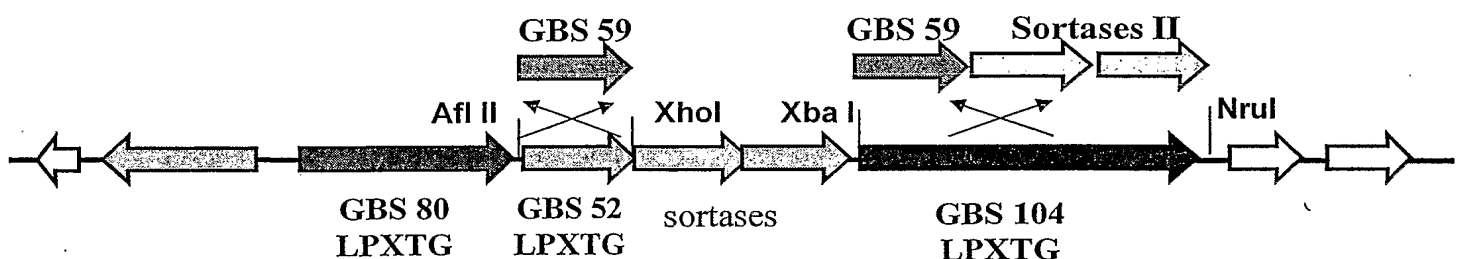


Oligo GBS67pAMXbafor AGTCAGTCTCTAGACGGCACAATAGGAGTTGTAAA

Oligo GBS67pAMBglrev CACCTGTCATAGATCTTAAGAATACTAAAGCGCATAA

2- Substitution of GBS52 or 104 with:

- GBS 59 alleles 515 or CJB
- GBS 59 allele CJB111 + sortases island II
- GBS 59 allele 515 + GBS 59 CJB111 + sortases island II



DETAILS:

a) Oligos to be used:

Oligo 59pAMAflfor1 AGTCAGTCCTTAAGCCGCATATTATTAATCATGTTG (allele 515)

Oligo 59pAMAflfor1 AGTCAGTCCTCGAGTTAACTTCCTCTGATTGACG (allele 515)

Oligo 59pAMAflfor2 AGTCAGTCCTTAAGAAGGAGTGGTGCTGCGGTAA (allele CJB111)

Oligo 59pAMXhorev2 AGTCAGTCCTCGAGTTAAGCTTCCTCTGATTGACG (allele CJB111)

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b) Oligos to be used:

Oligo GBS59XbaF CTAGTGATATATCTAGAGAAAAAG

Oligo Sort59NruR CTAGCTAGTCGCGACTTTTTCATTTTGGATTTCCCTTTC

Figure 234 B

3- Substitution of GBS104 with a fusion of GBS322-GBS67 to include GBS 322 into AI-1

- a) Construct 1: GBS67 complete sequence included
- b) Construct 2: Only part of GBS 67 was included (*deleted bold region*)

DETAILS:

a) Construct 1:

Legend:

Pink GBS322

Black GBS67

Black Bold: fragment of GBS67 eliminated in construct 2

Green PK motifs

Yellow E motifs

Red LPXTC

> gbs67-515 + 322

```

MRKYQKFSKILTLSLFCLSQIPLNTNVLGESTVPENGAKGKLVVKKTDQNKPLSKATFV
LKTTAHPESKIEKVTAELTGEATFDNLIPGDYTLSEETAPEGYKKTNQTWQVKVESNGKT
TIQNSGDKNSTIGQNQEELDKQYPPTGIYEDTKESYKLEHVKGSPNGKSEAKAVNPYSS
EGEHIREIPEGTLKRISVGDLAHNKYKIELTVSGKTIVKPVDKQKPLETDTTWTARTVSEV
KADLVKQDNKSSYTVKYGDTLSEAMSIDMNVLAKINNIADINLIYPETTLTVTYDQKSHTA
ISMKIETPATNAAGOTTATVDLKTNOQSVADOKVSLNTISEGIMTPEAATTIVSPMKTYSSAF
ALKSKEVLAEQQAQVSAQAANEQVSPAPVKSTSEVPAAKEEVKPTQTSVSQSTTVSPASV
AETPAPVAKVAPVRTVAAPRVASVKVVTPIKVEGASPEHVSAPAVPVTTTSPATDSKLOAT
EVKSPVPAQKAPTATPVAQPASTTNAVAHPENAGLQPHVAAYKERVASTYGVNEESTYRAC
DPGDHGKGLAYDFIVGTNOALGNKVAQYSTQNMANNISYVIWQOQKEYSN
INSYGPANTWINAMPDRGGVTANNDHVFHVSFNKDWVFVLDNSNS
MNNDGPNFQRHNKAKKAAEALGTAVKDILGANSNDRVALVTYGSDFDGRSVDVVKGFKE
DDKYYGLQTKFTIQTENYSHKQLTNNAEEIIRIPTEAPKAKWGSTTNGLTPEQQKEYYL
SKVGETFTMKAFMEADDILSQVNRNSQKIIVHVTGVPTRSYAINNFKLGAYESQFEQM
KKNGYLNKSNFLITDKPDDIKNGESYFLPLDSYQTQIISGNLQKLHYLDLNLNPKGI
IYRNGPVKEHGTPTKLYINSLKQKNYDIFNFGIDISGRQVYNEEYKKNQDGTGFKLKEE

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AFKLS DGEITELMRSFSSKPEYYTPIVTSADTSNNEILSKIQQQFETILTKENSIVNGTI
EDPMGDKINLQLGNGQILQPSDYTLQGNDGSVMKDG IATGGPNNDGGILKGVKLEYIGNK
LYVRGLNLGEGQKVTLTYDVKLDDSFISNKFYDTN GR TTL NPK SEDPNTLRDFPIPKIRD
VREYPTITIKNEKKLGEIEFIKVDKDNKKLLKGATFELQEFNEDYKLYLPIKNNNSKV
TGENGKISYKDLKDGKYQLIEAVSPEDYQKITNKPILTFEVVKGSIKNIIAVNKQISEYH
EEGDKHLITNTHIPPKGI KICU KGILSFILIGGAMMSIAGGIYWKRYKKSSDMSIKK
D

Figure 234 C

b) Construct 2:

>gbs67-515 deleted+ 322

MRKYQKFSKILTLSLFCLSQIPLNTNVLGESTVPENGAKGKLVVKKTDQNKPLSKATFV
 LKTTAHPESKIEKVTAELTGEATFDNLIPGDYTLSEETAPEGYKKTNQWQVKVESNGKT
 TIQNSGDKNSTIGQNQEELDKQYPPTGIYEDTKESYKLEHVKGSPNGKSEAKAVNPYS
 SEGEHIREIPEGTLKRISSEVGDLAHNKYKIELTVSGKTIVKPVDPKQKPLETDTTW
 TARTVSEVKADLVKQDNKSSYTVKYGDTLSVISEAMSIDMNVLAKINNIADINLIYPETTLTV
 TYDQKSHATATSMKIETPATNAAGQTTATVDLKTNGVSVADQKVSLENTISEGMTPEAATT
 VSPMKTYSSAPALKSKEVLAQEQAVSQAAANEQVAPVKSITSEVPAAKEEVKPTQTS
 VSOSTTVGPAASVAAETPAPVAKMAPVRTVAAPRVASVKVYTPKVVETGASPEHVSAPAVE
 VTTTSPATDSKLQATEVKSVPVAQKAPTATPVAOPASTTNAVAHPENAGLOPHVAAYK
 EKVASTYGVNEPSTYRAGDPGDHCKGLAVDFMGTNGALGNKYAOYSTONMAANNISY
 WQQKFYSNTNSIYGPANTWNAMPDRGGVTANHMDHVHVSFNKGESYFLPLDSYQQTQ
 IISGNLQKLHYLDLNLNYPKGTIYRNGPVKEHGTPTKLYINSLKQKNYDIFNFGIDISGRQ
 VYNEEYKKNQDGTGFKLKEEAFKLSDEITELMRSFSSKPEYYTPIVTSADTSNNEILSKI
 QQQFETILTKENSIVNGTIEDPMGDKINLQLGNGQILQPSDYTLQGNDGSVMKDGATGG
 PNNDGGILKGVKLEYIGNKLYVRGLNLGEGQKVTLTYDVKLDDSFISNKFYDTNGRTTLN
 PKSEDPNTLRDFPIPKIRDVREYPTITIKNEKKLGEIEFIKVDKDNKLLKLGATFELQEFNE
 DYKLYLPIKNNNSKVVTGENGKISYKDLKDGKYQLIEAVSPEDYQKITNKPILTFEVVKG
 IKNIIAVNKQISEYHEEGDKHLITNTHIPPKGIPKIGKILSFILIGGAMMSIAGGIYIWKRY
 KKSSDMSIKKD

Oligos to be used:**Oligo GBS67pAMXbafor (vedi operone)**

AGTCAGTCTCTAGACGGCACAAATAGGAGTTGTAA

XbaI

Oligo GBS67soe1rev

GCAACGCGGATATGCTTTC TAACGGCTTTTGTGTCCACT

Oligo GBS322soe2for

GACAAACAAAAGCCGTTAATAACAGATACCAAGGTGGACAG

Oligo GBS322soe2rev1 (per costruito non delete in 67)

GAGTACGAAGACAACATCTTGTAAATGATACGTCGACG

Oligo GBS322soe2rev2 (per costruito delete in 67)

TAAAAAGTAACTCTCCCCCTTGTAAATGATACGTCGACG

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Oligo fine67soe3for1 (per costruito non deleto in 67)

GACGTATCATTAACAAAGATGTTGTCTTCGTACTCGAT

Oligo fine67soe3for2 (per costruito non deleto in 67)

GACGTATCATTAACAAAGGGGAGAGTTACTTTTTATTTC

Oligo GBS67pAMBglrev (vedi operone)

CACCTGTCATAGATCTTAAGAATAC TAAAGCCGCA TAA

BglII

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Figure 234 D

PCR Soe1: GBS67pAMXbafor + GBS67soe1rev 727 bp

PCR Soe2 non del: GBS322soe2for + GBS322soe2rev1 1260 bp

PCR Soe2 del: GBS322soe2for + GBS322soe2rev2 1260 bp

PCR Soe3 non del: fine67soe3for1 + GBS67pAMBglrev 2061 bp

PCR Soe3 del: fine67soe3for2 + GBS67pAMBglrev 1419 bp

PCR Soe4 non del. PCR25: GBS67pAMXbafor + GBS67pAMBglrev 4000 bp

Substrato PCRSoe1, 2, 3 non del

PCR Soe4 del, PCR26: GBS67pAMXbafor + GBS67pAMBglrev 3312 bp

Substrato PCRSoe1, 2, 3 del

4- Substitution of GBS 52 with a fusion of GBS322-GBS52 to include GBS 322 into AI-1

(same legend as for GBS67 derivatives)

a) Construct 1: GBS52 complete sequence included

b) Construct 2: Only part of GBS 52 was included (*deleted bold region*)**DETAILS:**

a) Construct 1:

>GBS322-52 senza delezione di 52 (B) PCR 24

MKMNKKVLLTSTMAASLLSVASVQAQETDTLWTARTVSEVKADLVKODNK
 SSYTVKYGDILLSVISEAMSIDMNVLAKEINNIADINLIYPETTLTVTYDQK
 SHATSMKIETPATNAAGQTTATVDLKTNOVSVADQKVSINTISEGNTRE
 AATTIVSPMKTYSSAPALKSKEVLAQEQAVSQAAANEQVSPA
 RVKSITSEVPAAKEELAKPTQTSYSQSTTVSPASVAAETPAPVAKVAPVRTVAAPRVAS
 MKVVTPEKVTGASPEHVSAPAVPVTTTSPATDSKLOATEVKSPVPAQKAF
 IATPVAQPASTTNAAHPENAGLOPHVAAYKEKVASTYGVNEFSTYRAG
 DPGDHGKGLAVDFIVGTNOALGNKVAQYSTQNMANNISYIMWQOKEYSN
 INSIYGPANTWNAMPDRCCVITANHYDHVHVSFNK HQLTIVHLEARDIDRPNPQL
 EIAPKEGTPIEGVL YQLYQLKSTEDGDLLAHWNSLTITELKKQAQQVFEA
 TTNQQGKATFNQLPDGIYYGLAVKAGEKNRNVSAFLVDLSEDKVIYPKII
 WSTGELDLLVGVVDGDTKKPLAGVVFELYEKNRTPIRVKNGVHSQDIDA
 AKHLETDSSGHIRISGLIHGDYVLKEIETQSGYQIGQAETAVTIEKSKTV

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TVTIENTKKVPTPKVPSRGGLEKQQAMALVIIGGILIALALRLLSKH
RKHQNKD

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Figure 234 E

b) Construct 2:

>GBS322-52 (A) PCR 23

MKMNNKKVLLTSTMAASLLSVASVQAQETDTTWTARTVSEVKADLVKQDNK
 SSYTVKYGDTLSVISEAMSIDMNVLAKEINNIADINLIYPETTLTVTYDOK
 SHTATSMKIEPTPATNAAGOTTATVDLKTNOVSVADOKVSLNTISEGMTPE
 ATTIVSPMKTYSSAPALKSKEVLAQEQAVSQAAAANEQVSPA
 PVKSITSEVPAAKEEVKPTQTSVSOSTTVSPASVAAETPAPVAKVAPVRTVAAPRVAS
 VKVYTPKMETGASPEIIVSAPAVPVTTTSPATDSKLOATEVKSVPAQKAF
 IATPVAQPASTTNAAVAHPENAGLOPHVAAVKEKVASTYGVNEFSTYRAG
 DPGDHGGKGLAVDEIVGTNQAQGNKVAQYSTONMAANNISYVWQCKEYSN
 NSLYGPANTWNAMPDRGGVTANHVDHVHVSNNK
 QGKATFNQLPDGIYYGLAVKAGEKNRNVSAFLVDLSEDKVIYPKII
 WSTGELDLLKVGVDGDTKKPLAGVVFELYEKNRTPIRVKNGVHSQDIDA
 AKHLETDSSGHIRISGLIHGDYVLKEIETQSGYQIGQAETAVTIEKSKTV
 TVTIENKKVPTPKVPSRGGLEKICEQQAMALVIIGGILIALRLLSKH
 RKHQNKD

Oligos to be used:

Oligo 322Aflfor1

AGTCAGTCCTTAAGGATATTATAGTCTCGGACTA

Afl II

Oligo 52 soe1 forA

CAAGGAAAGGCTACATTTAACG

Oligo 52 soe1 forB

CATCAGTTGACGATTGTTTCATC

Oligo52 soe1revA

AAATGTAGCCTTTCCTTGTTTGTTAAATGATACGTCGACG

Oligo52 soe1revB

AACAATCGTCAACTGATGTTTGTTAAATGATACGTCGACG

Oligo 52Xhorev

AAGACCTCCTCGAGATGGCACTT

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Xho I

PCR Soe1A: Oligo 322Aflfor1+ Oligo 52 soe1 revA 1370 bp

PCR Soe2A: Oligo52 soe1forA + Oligo 52Xhorev 520 bp

PCR Soe3A: Oligo 322Aflfor1 + Oligo 52Xhorev 1846 bp (con PCR Soe1A + PCR Soe2A)
(PCR23)

PCR Soe1B: Oligo 322Aflfor1+ Oligo 52 soe1 revB 1370 bp

PCR Soe2B: Oligo52 soe2forB + Oligo 52Xhorev 742 bp

PCR Soe3B: Oligo 322Aflfor1 + Oligo 52Xhorev 2068 bp (con PCR Soe1B + PCR Soe2B)
(PCR 24)

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Figure 235

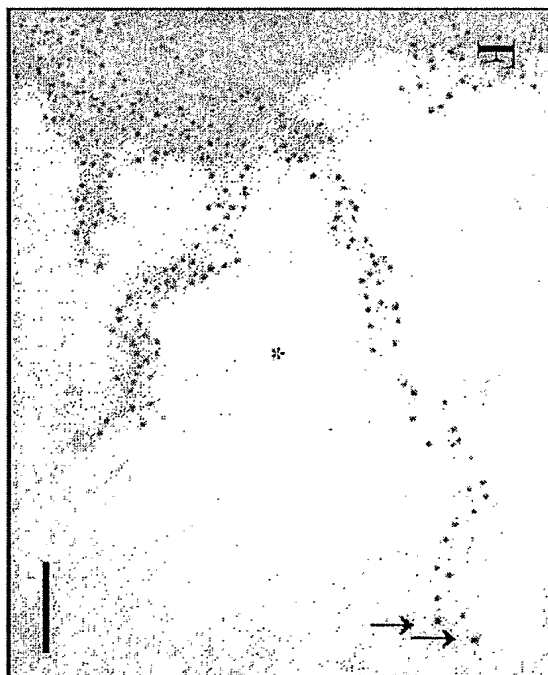


Figure 236



Strain variability - GBS67: 2 alleles

1 MRKYQKPSKILITLSLFCISQIPLNTNVLGSEVPENGAKGLVVKTTDDQ 50
1NVLGESTVPENGAKGLVVKTTDDQ 25
51 NKPLSKATFVLKTAHPESKIEKVTAELTGEATFDNLIPGDYTLSEETAP 100
26 NKPLSKATFVLKTPSHSESKEVKVTEVTGEATFDNLTPGDYTLSEETAP 75
101 EGYKKTQWQVKNVSGKTTIQNSGDKNSTIGNQBELDKQYPTGIYE 150
76 EGYKKTQWQVKNVSGKTTIQNSDDKKSIIIEQRQBELDKQYPLTCAYE 125
151 DTKEYSKLEHVKGSPVNGKSEAKVNPYSSEGEHIREIPEGTLSKRISV 200
126 DTKEYSNLEHVKNISPGKLEAKVNPYSSEGEHIREIQEGLTSKRISV 175
201 GDLAHNKYKIELTVSGKTIKVPDKQKPLDVVFLDNSNSMNDGNPFOR 250
176 NDLDHNKYKIELTVSGKSIKTIKNDKDEPLDVVFLDNSNSMKNKGN... 222
251 HNKAKAAEALGTAVKDIILGANSNDRVALVTYGSDFDGRSVDVVKGFKE 300
223 .NKAKKAGEAVENTIKDVLGANVENRAALVTYGSDFDGRVTVKVGFE 271
301 DDKYGLQTKFTIQTENYSHKQLTNNABEILKRIPEAPKAKWGSTNGL 350
272 .DPYYGLETSFTVQTDNDYSYKFTNTAADIIKKIPKEAPEAKWGTSIGL 320
351 TPEQKRYLLSKVGEFTMKAFWEADDDILSQVNRNSQKLIHVHVDGVPT 400
321 TPEKRYDLSKVGEFTMKAFWEADTILSSIQRSKRLIVHLTDGVPT 370
401 SYAINNFKLGASVESFEQKKNNGYLNKSNFLIDKPEDIKNGESYFLF 450
371 SYAINSFVKGSTVANOFEIKGKGYLDKNVFLTDDEPKIKNGESYFLF 420

451 PLDSYQTIISGNLQKLHYLDLNLNYPKGTIYRNGPVEHGTPTKLYINS 500
421 PLDSYQTIISGNLQKLHYLDLNLNYPKGTIYRNGPVEHGTPTKLYINS 480
501 LKQKNYDIFNFGIDISGRQVYNEEYKKNQDGTFOKLKEAFKLSDGEIT 550
471 LKQKNYDIFNFGIDISGRQVYNEEYKKNQDGTFOKLKEAFKLSDGEIT 520
551 ELMRSFSKPEYYTPIVTSADTSNNELLSKIQQQFETILTKEINSVNGTI 600
521 ELMRSFSKPEYYTPIVTSADTSNNELLSKIQQQFETILTKEINSVNGTI 580
601 EDPMGDKINLQNGQTLQPSDYTLQNGDGSVMKDGDIATGSPNNDGGILK 650
571 EDPMGDKINLHNGQTLQPSDYTLQNGDGSIMKDSIATGSPNNDGGILK 620
651 GVKLEYIGNKLYVRGLNGLGEGQKVTLYDVKLDSDSFISNKFYDTNGRTTL 700
621 GVKLEYIKNKLYVRGLNGLGEGQKVTLYDVKLDSDSFISNKFYDTNGRTTL 670
701 NPKSEDENTLRDFPIPKIRDVREYPTITIKNEKKGIEFTIKYDKONNKL 750
671 NPKSEEDTLRDFPIPKIRDVREYPTITIKNEKKGIEFTIKYDKONNKL 720
751 LLKGATFELQEFNEDYKLYLPIKNNNSKVVTGENGKISYKDLKDGKYQLI 800
721 LLKGATFELQEFNEDYKLYLPIKNNNSKVVTGENGKISYKDLKDGKYQLI 770
801 EAVSPEDYQKITNKPILTTEVVKSGSIKNIIVANKQISEYHEEGDKHLITN 850
771 EAVSPDYQKITNKPILTTEVVKSGSIQNIIVANKQISEYHEEGDKHLITN 820
851 THIPPKGIIPMTGGKILSFILIGAMMSIAGGIYIWKRYKKSSDMSIKK 900
821 THIPPKGI..... 828

Differences
between strains
2603 and H36B
(AA not matching/AA
total and % of homology)

114 / 828 (87,1%)

Figure 237

Strain variability - GBS67 Allele I (2603)

Strain	Differences in comparison with 2603 (% of homology)
2603	-
18RS21	1/833 (99.9%)
CJB111	14/833 (98.3%)
515	2/833(99.8%)

Figure 238

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Strain variability - GBS67 Allele II (H36b)

Strain	Differences in comparison with H36b (% of homology)	FACS (α -67 from 2603)
H36B	-	444
1169	10/823 (98.8%)	443
090	9/316 Stop codon (8G to 7G)	0
CJB110	11/824 (98.7%)	245

Figure 239